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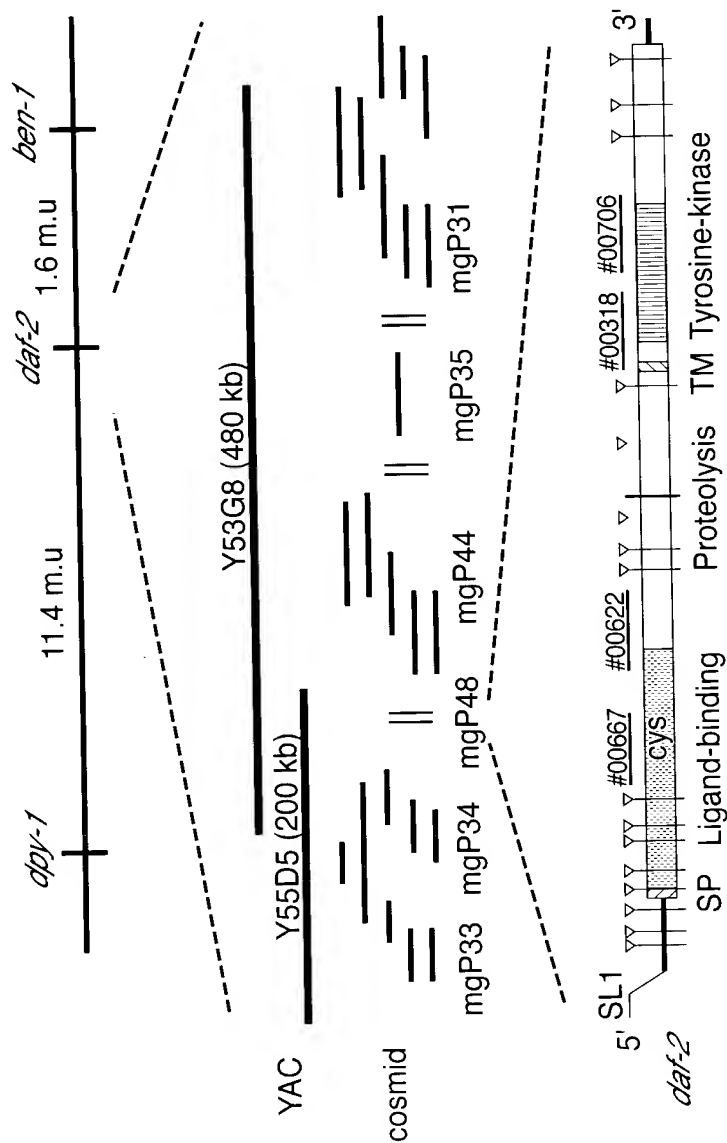


Fig. 1



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121 KIFPNLRVIGGRSLIQHYALIIYRNPDL EIGLDKLSVIRNGGVRIIDNRKLCYTKTIDWK
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241 SNTTCQKSCAYDRLLPTKEIGPGCDANGDRCHDQCVGGCERVNDATAACHACKNVYHKGKC
301 IEKCDAHLYLLLQRRCVTREQCLQLNPVLSNKTVP I KATAGLCSDKCPDGYQINPDDHRE
361 CRKCVGKCEIVCEINHVIDTFPKAQAIRLCNIIDGNLTIEIRGKQDSGMASELKDIFANI
421 HTITGYLLVRQSSPFISLNMFRNLRRIEAKSLFRNLYAITVFENPNLKKLFDSTDTLD
481 RGTVSIANNKMLCFKYIKQLMSKLNIPLDPIDQSEGTNGEKAICEDMAINVSITAVNADS
541 VFFSWPSFNITDIDQRKFLGYELFFKEVPRIDENMTIEEDRSACVDSWQSVFKQYYETSN
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661 LAQVDSDAIHITWEAPLQPNGDLTHYTIMWRENEVSPYEEAEKFCTDASTPANRQRTKDP
721 KETIVADKVPDIPSSRTVAPTLTMMGHEDQOKTCAATPGCCSCSAIEESSEQNKKKRPD
781 PMSAIESSAFENKLLDEVLMPRDTMRVRRSIEDANRVSEELEKAENLGKAPKTLGGKKPL
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901 LDNSYVIRNLKHYTLYAISLSACQNMTVPGASCSISHRAGALKRTHITDIDKVLNETIE
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1021 QNLADGRYFVSVTATSVHGAGPEAESSDPIVVMTPGFFTVEIILGMLLVFLILMSIAGCI
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1201 PAMVVMEMMDLGNLRDYLRSKREDEVFNEDCNFFDIIPRDKFHEWAAQICDGMAYLESL
1261 KFCHRDLAARNCMINRDETVKIGDFGMARDLFOHDQKPSGKRMPVRWMSPELKD GKF
1321 DSKSDVWSFGVVLYEMVTLGAQPYIGLSNDEV LNYIGMARKVIKKPECCENYWKVMKMC
1381 WRYSRDRPTFLQLVHLAAEASPEFRDLSFVLTDNQMILDDSEALDLDIDDTDMNDQV
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1501 DEEYALMNHSGGPSDAEVRTYAGDGDVERDVRENDVPTRRNTGASTSSYTGGGPYCLTN
1561 RGGSNERGAGFGEAVRLTDGVGSGHLNDDDDVEKEISSMDTRRSTGASSSSYGVPTNWS
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1681 TEPKNYRNNGSPSRNGNSRDIFNGRSAFGENEHLIEDNEHHPLV

Fig. 2A



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Fig. 2B-1



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Fig. 2B-2



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Fig. 2B-3

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 Inr RGSVRIEKNELCYLATIDWSRLIDSVEDNYIVLNKDDNE.EGG.DICPGTAKGKTNCPATVINGQFVER.....CWTHSHCCQKVC.....PTIC
 Dinr RGVRIEKNHKLCDRTIDWLELIDAEENESQVLTENGKEKESLSKCPGEIRIEEGHNTATEGELNASCOLHNRRRLCWNKSKLCCQTKC.....PEKC
 DAF-2 NGGVRIIDNRKLCYTKTIDWKHLITSSINDVVDNAEYAVTETGLMCPRGACEEDKESKCHYLEEKNQEGQVERVQSCWSNNTTCQKSCAYDRLLPTKE

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 Inr KSHGCTAEGLCCHSECLGNC SQDDPTKCVACRNFIDRCVETCPPPYVHEQDWKCVNFSFCQDPHHKCKNSRRQCHQYVIHNNKCIPCEPSGYTMN
 Dinr RNNCIDHTCCSQDCLGGGVIDKNGNESCSIRNVSFNNICMDSCKPGYVQF.DSRCVTANECITITKFTNSVYSG...IPYNGCQITHCPTGY.OK
 DAF-2 IGPGDANGDRCHDQCVEGC ERVNDATACHACKNVYHKKCKIEKCD AHLALALQRCVTRCQCLQNPVLSNKTVP....IKATACGLGSDKCPDGYQIN
 Y (mg43)

IGF-IR GSQSMYICIPCEGCPKVCIEEEKKTKTIDSVTSQMLQGCTIFKGN..LLENRR..GNNIASETENFMGLTEVATGYVKIRHSHARVSLSEKKNRLRLIG
 Inr SSN.LLCTPCLEGPCPKVCHLLEGEKTIIDSVTSQELRGCTWINGS..LLENRRG..GNNIAAET.EANIGLIEEISGYLKIERSYALVSLSEFFRKLRLIRG
 Dinr SENKRMCEPCPGG...KDKCECSSLIDSLEAREFHCITITGTEP...LSLRSCAHVMDDELKYGLAAWHKHQSSLMVHLTYGKSIKKEFQSLTEISG
 DAF-2 PDDHRECRKCVCKEIVGEI...NHVIDMFPFAQAIRLCNIIDEN..LLETRGKQDSGMASETKIDIFANTHTITGYLLVQSSPFIISINMERNLRRHEA
 L (mg43)

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 Inr ETLLEIGN.VSFVLDNQLQQLWDWSKNEHTITQGLFFHYNPKLCGLSEIHKMEEVSGTKGR.QERNIDIALKTNGDQASCENELKFSYIRTSFD.....
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 DAF-2 KSLFR.NLVAITVLENPNLKKLLE.D.STTDTLDRGTVSIANNKMLGFKYIKQLMSKLNIP...LDPILOQSEGTNGEKACEDMANVNSITAVNADS....
 L (e1368) T (e1365)

E S
 L (sa187)
 R
 N (sa229)

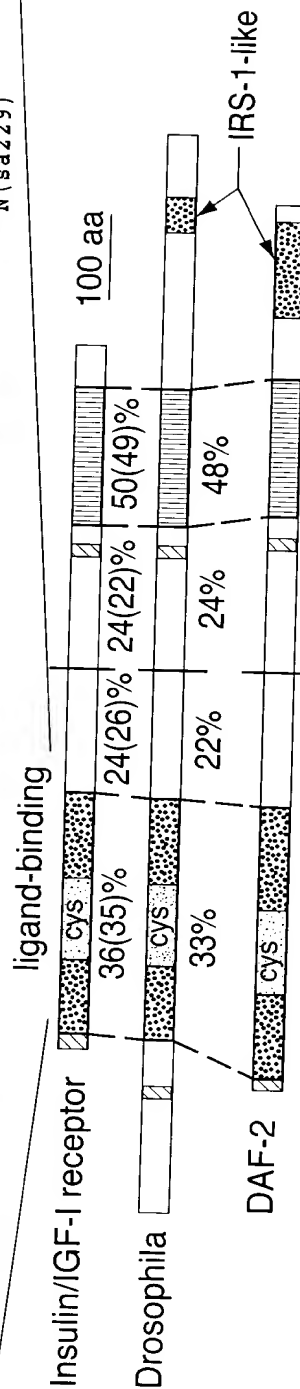


Fig. 2C-1

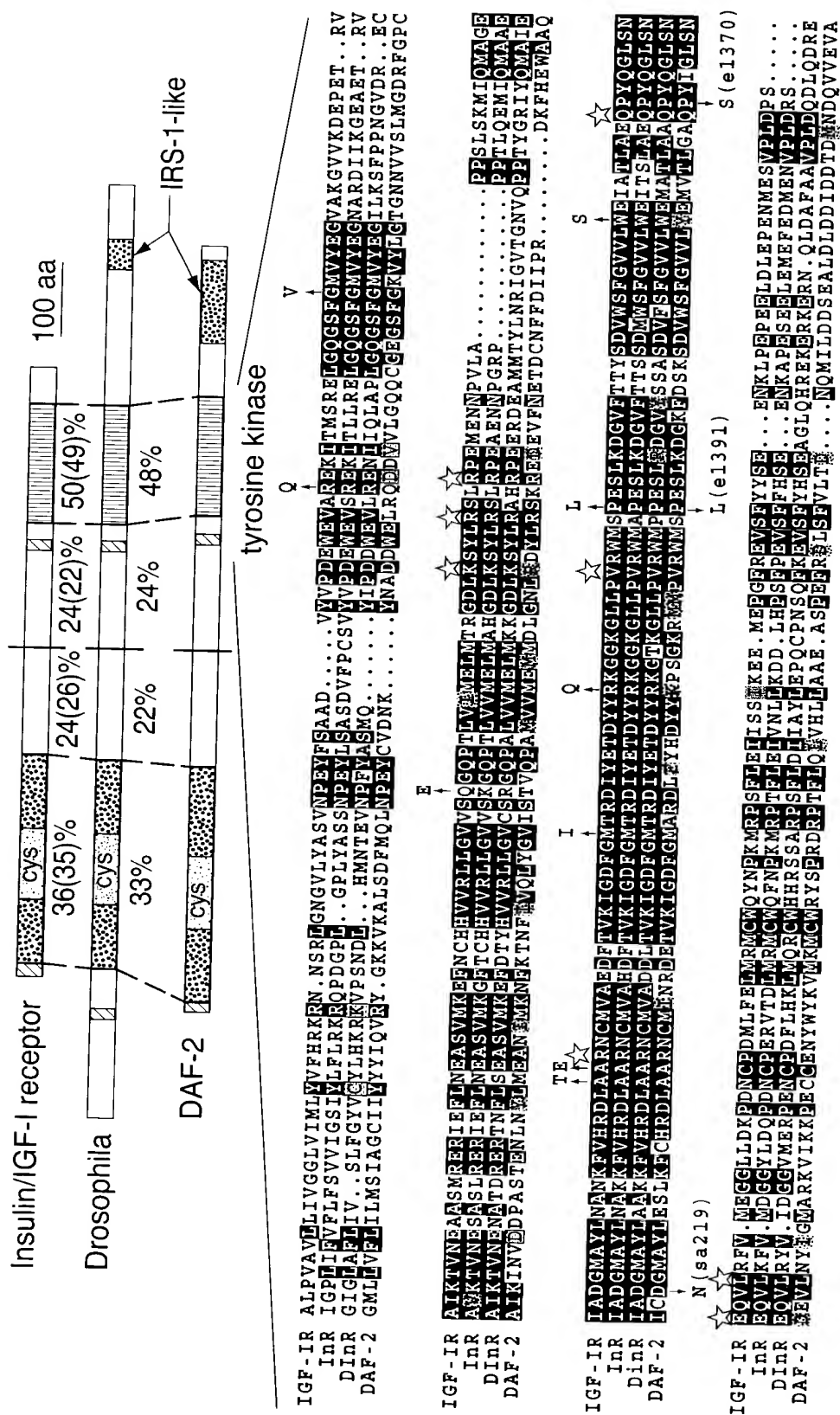


Fig. 2C-2

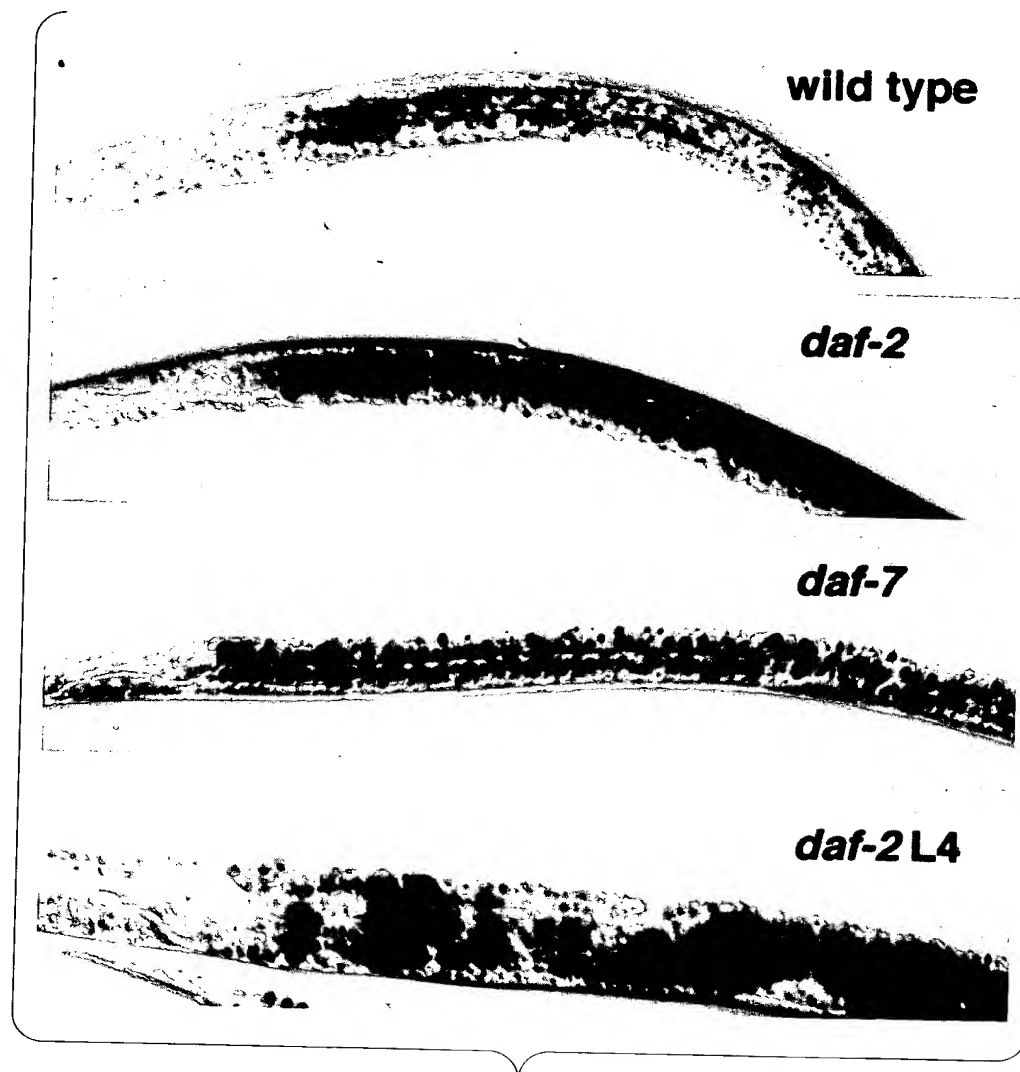


Fig. 3

Fig. 4

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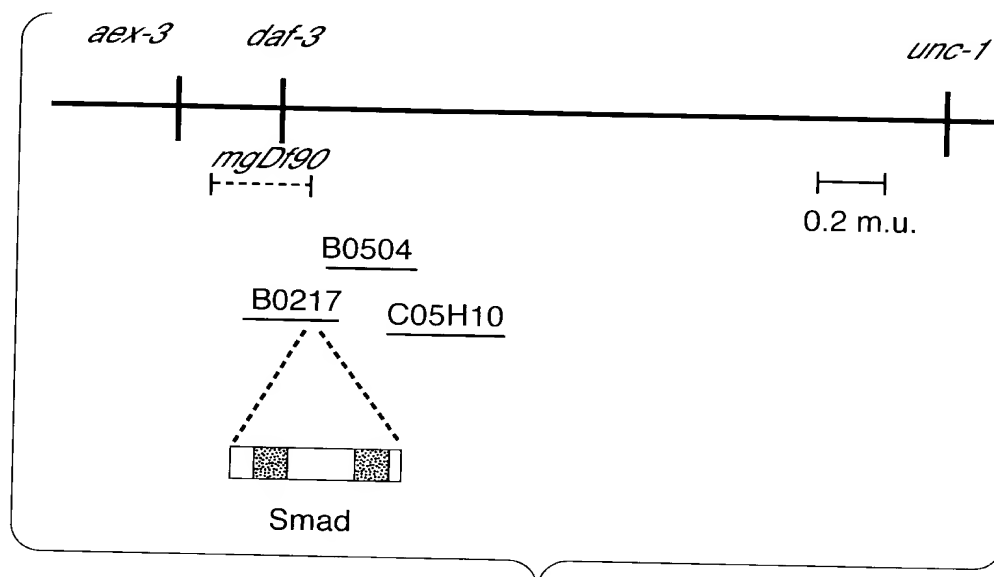


Fig. 5A

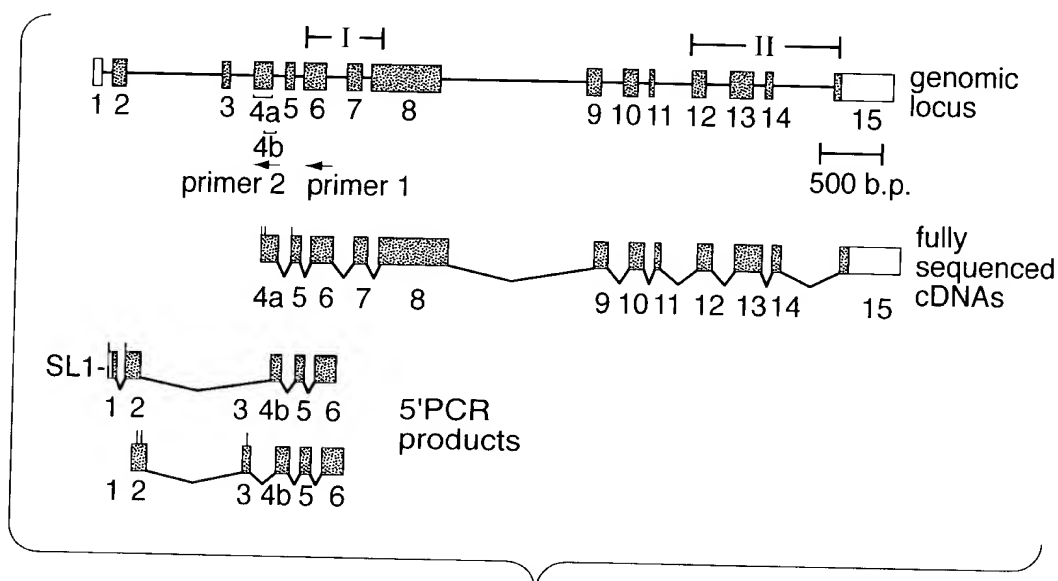


Fig. 5B

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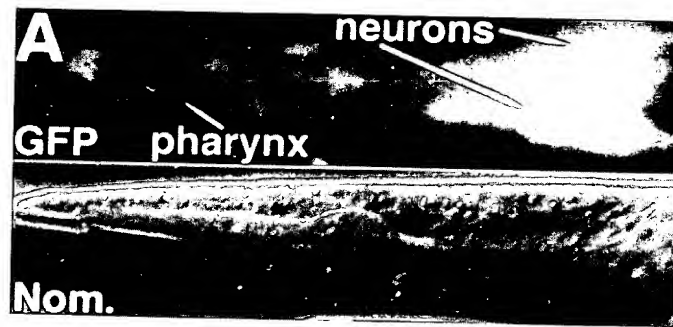


Fig. 6A

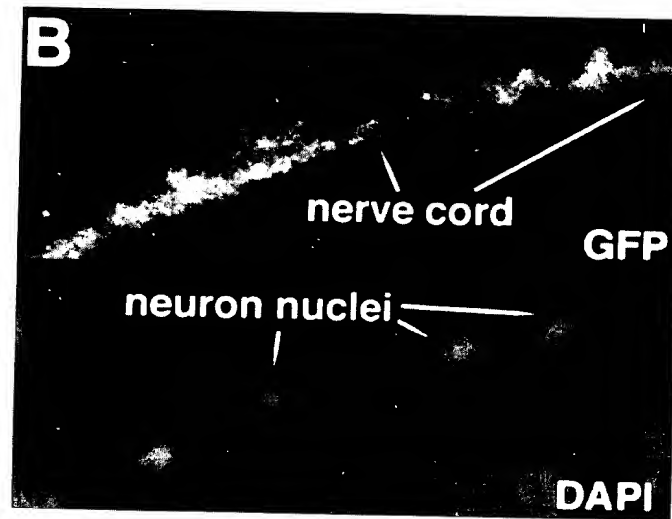


Fig. 6B

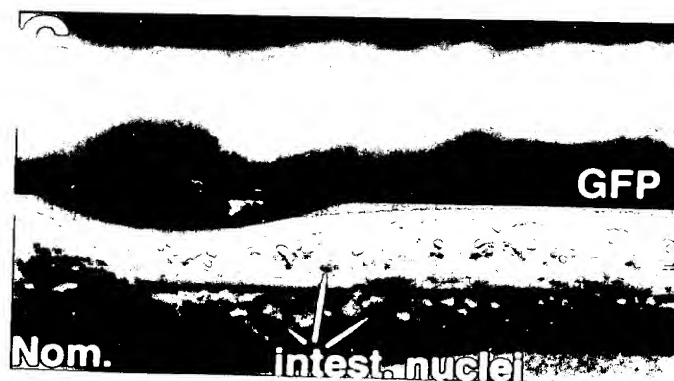


Fig. 6C

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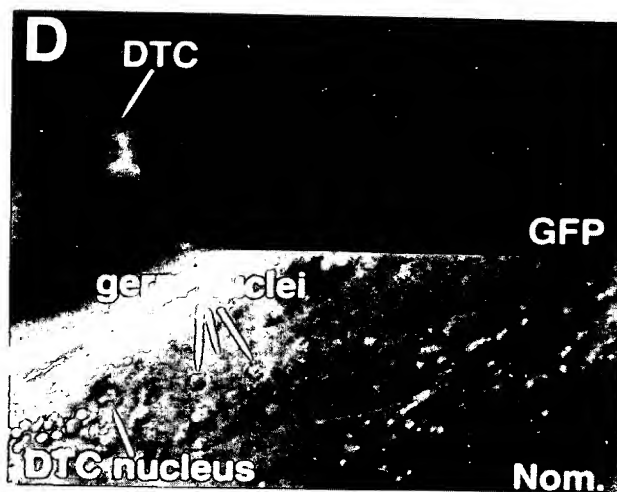


Fig. 6D

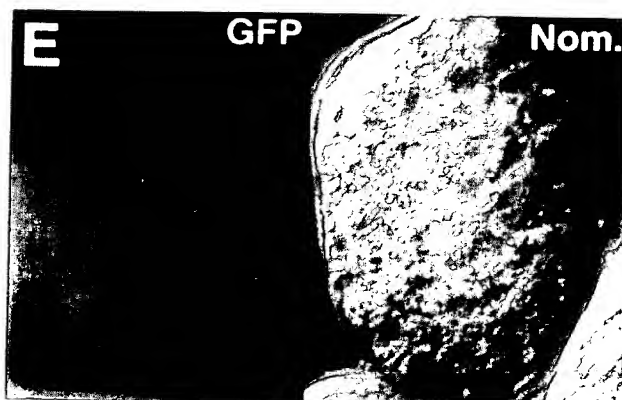


Fig. 6E

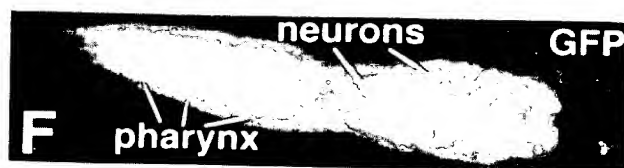


Fig. 6F



Fig. 6G

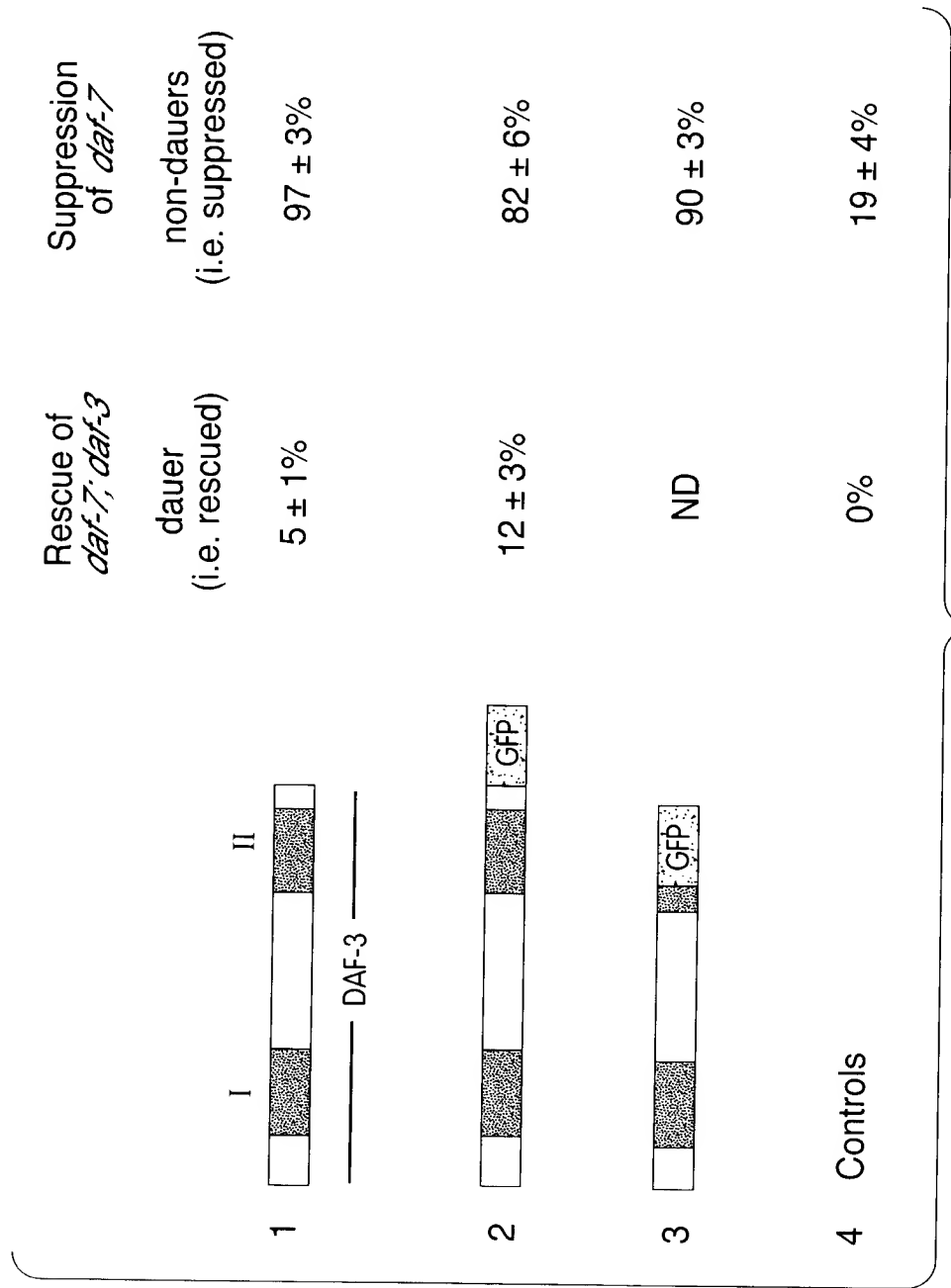


Fig. 7

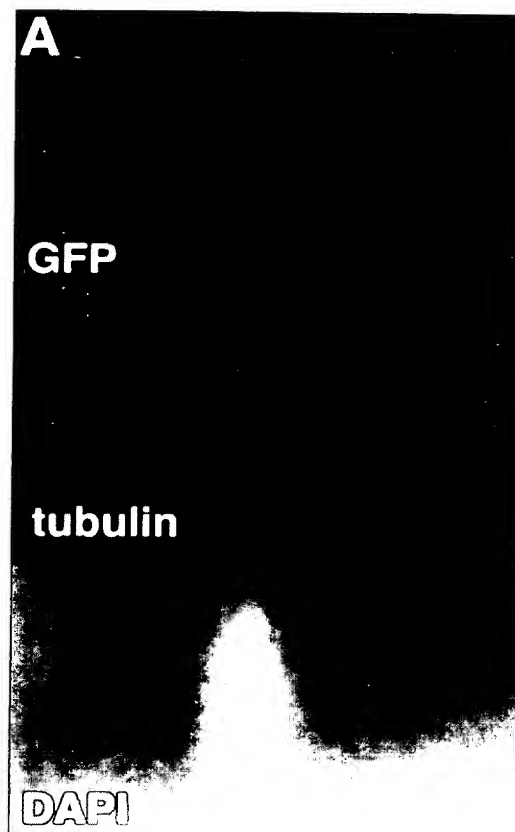


Fig. 8A

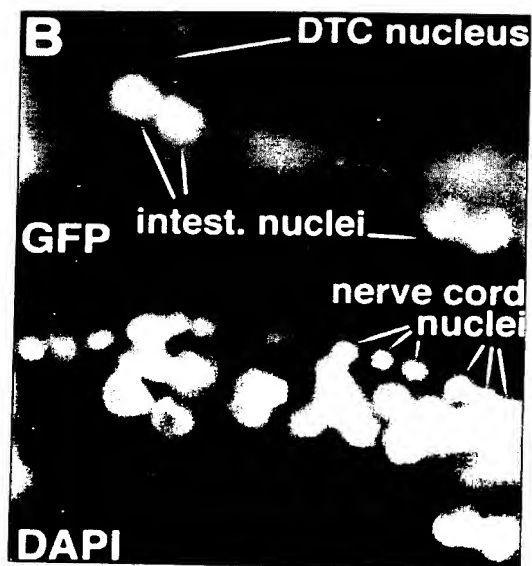


Fig. 8B

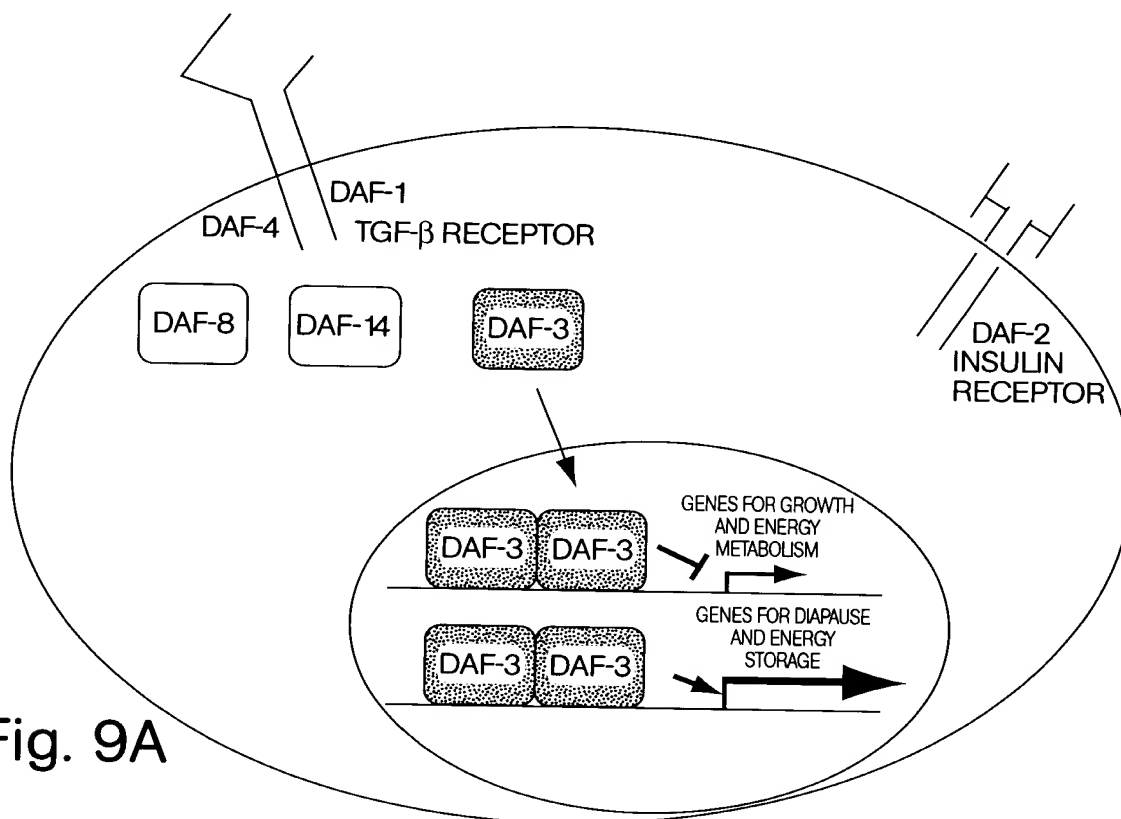


Fig. 9A

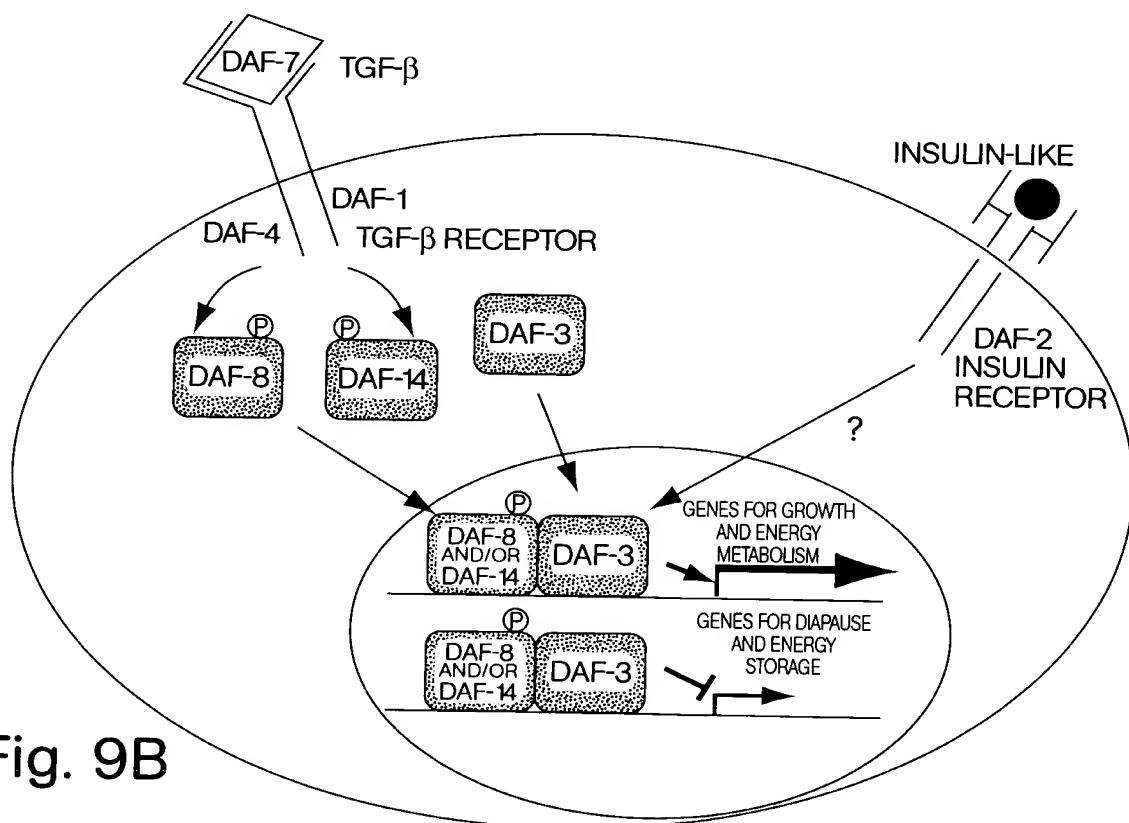


Fig. 9B



Diagram illustrating the DAF-12 signaling pathway in *C. elegans*, showing the progression from pheromone to dauer stage, and the role of various genes in this process.

Genes and their interactions:

- Sensory neuron ASJ:** Contains genes *daf-11*, *daf-21*, *che-2*, *osm-3*, etc. This neuron is activated by pheromone and promotes the dauer stage.
- Sensory neurons ASI, ASG, ADF:** Contains genes *daf-1*, *daf-4*, *daf-7*, *daf-8*, *daf-14*, *daf-3*, *daf-5*. These neurons are activated by the ASJ neuron and promote the dauer stage.
- DAF-12:** A transcription factor that promotes the dauer stage. Mutations in *daf-12* lead to the L3 stage.
- Other genes:**
 - daf-16*: Promotes the dauer stage.
 - daf-2* and *age-1*: Inhibit the dauer stage.
 - daf-22* and *daf-6*: Promote the dauer stage.

Phenotypes:

- Daf-d:** dauer stage.
- Daf-c:** dauer stage.
- Daf-d:** dauer stage.
- Daf-c:** dauer stage.
- Daf-d:** dauer stage.
- Daf-d:** dauer stage.



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1	atgaagctaa	tagcaacttc	tcttctagtt	cccgacgagc	acacacccgat
51	gatgtcacca	gtgaatacaa	ctacaaagat	tctacaacgg	agtgggtatta
101	aaatggaaat	cccgccatat	ttggatccag	acagtcagga	tgatgacccg
151	gaagatggtg	tcaactaccc	ggatccagat	ttatttgaca	caaaaaacac
201	aaatatgacc	gagtacgatt	tggatgtggt	gaagcttgga	aaaccagcag
251	tagatgaagc	acggaaaaag	atcgaagttc	ccgacgctag	tgcgcccgcca
301	aacaaaattg	tagaatattt	gatgtattat	agaacgttaa	aagaaagtga
351	actcatacaa	ctgaatgcgt	atcggacaaa	acgaaatcga	ttatcgttga
401	acttggtcaa	aaacaatatt	gatcgagagt	tcgaccaaaa	agcttgcgag
451	tccctgggtga	aaaaattgaa	ggataagaag	aatgatctcc	agaacctgat
501	tgatgtgggt	ctttcaaaag	gtacaaaata	taccggttgc	attacaattc
551	caaggacact	tgatggccgg	ttacaggtcc	acggaagaaa	aggtttccct
601	cacgtagtct	atggcaaact	gtggagggtt	aatgaaatga	caaaaaacga
651	aacgcgtcat	gtggaccact	gcaagcacgc	atttgaaatg	aaaagtgaca
701	tgggtatgct	gaatccctat	cactacgaaa	ttgtcattgg	aactatgatt
751	gttgggcaga	gggatcatga	caatcgagat	atgccgccgc	cacatcaacg
801	ctaccacact	ccaggtcggc	aggatccagt	tgacgatatg	agtagattta
851	taccaccagc	ttccattcgt	ccgcctccga	tgaacatgca	cacaaggcct
901	cagcctatgc	ctcaacaatt	gccttcagtt	ggcgcaacgt	ttgcccatcc
951	tctcccacat	caggcgccac	ataaccagg	ggtttcacat	ccgtactcca
1001	ttgctccaca	gacccattac	ccgttgaaca	tgaacccaat	tcgcgaaatg
1051	ccgcaaatgc	cacaaatgcc	accacctctc	catcagggat	atggaatgaa
1101	tgggcccag	tgctcttcag	aaaacaacaa	tccattccac	caaaatcacc
1151	attataatga	tattagccat	ccaaatcact	attcctacga	ctgtgggtccg
1201	aacttgtacg	ggttttccaac	tccttatccg	gatttttcacc	atcctttcaa
1251	tcagcaacca	caccagccgc	cacaactatc	acaaaaccat	acgtcccaac
1301	aaggcagtca	tcaaccagg	caccaaggtc	aggtaccgaa	tgatccacca
1351	atttcaagac	cagtgttaca	accatcaaca	gtcaccttgg	acgtgttccg
1401	tcggtactgt	agacagacat	ttggaaatcg	atTTTTtgaa	ggagaaagtg
1451	aacaatccgg	cgcaataatt	cggtctagta	acaaattcat	tgaagaattt
1501	gattcgccga	tttgtggtgt	gacagttggt	cgaccgcgga	tgacagacgg
1551	tgaggttttg	gagaacatca	tgccggaaga	tgcaccatat	catgacattt
1601	gcaagttcat	tttgaggctc	acatcagaaa	gtgtaacttt	ctcaggagag
1651	gggcccaga	ttagtgattt	gaacgaaaaa	tggggaacaa	ttgtgtacta
1701	tgagaaaaat	ttgcaaattg	gcgagaaaaa	atgttcgaga	ggaaatttcc
1751	acgtggatgg	cggattcatt	tgctctgaga	atcgttacag	tctcggactt
1801	gagccaaatc	caattagaga	accagtggcg	tttaaagttc	gtaaagcaat
1851	agtggatgga	attcgctttt	cctacaaaaa	agacgggagt	gtttggcttc
1901	aaaaccgc	gaagtacc	gtatttgtca	cttctgggta	tctcgacgag
1951	caatcaggag	gcctaaagaa	ggataaagt	cacaaagttt	acggatgtgc
2001	gtctatcaaa	acgtttggct	tcaacgtttc	caaacaaatc	atcagagacg
2051	cgcttctttc	caagcaaattg	gcaacaatgt	acttgcaagg	aaaattgact

Fig. 11A-1



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2101	ccgatgaatt	atatctacga	gaagaagact	caggaagagc	tgccaaggga
2151	agcaacacgc	accactgatt	cattggccaa	gtactgttgt	gtccgtgtct
2201	cgttctgcaa	aggatttgga	gaagcatacc	cagaacgccc	gtcaattcat
2251	gattgtccag	tttggattga	gttgaaaatc	aacattgcct	acgatttcat
2301	ggattcaatc	tgccagtaca	taaccaactg	cttcgagccg	ctaggaatgg
2351	aagattttgc	aaaattggga	atcaacgtca	gtgatgacta	aatgataact
2401	tttttcactc	accctactag	atactgattt	agtcttattc	caaateatcc
2451	aacgatatac	aactttttcc	tttgaacttt	gcatactatg	ttatcacaag
2501	ttccaagcag	tttcaataca	aacataggat	atgttaacaa	cttttgataa
2551	gaatcaagtt	accaactggt	cattgtgagc	tttgagctgt	atagaaggac
2601	aatgtatccc	atacctcaat	ctttaatagt	catcagtcac	tggtcccgcg
2651	ccaatttttt	cgatttcgcat	atgtcatata	ttgcaccgtg	gcccttttta
2701	ttgtaacttt	taatataatt	tcttcccaac	ttgtgaatat	gattgatgaa
2751	ccaccatttt	gagtaataaa	tgtatttttt	gtgg	

Fig. 11A-2



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1	gtaatcaa	at	tgtaaagg	aa	aatatta	at	agtcagag	ta	cacataa	aatg
51	ggtgatc	atc	ataattt	aac	gggcctt	ccc	ggtacct	cca	tcccgcc	caca
101	gttcaac	tat	tctcagc	cccg	gtaccag	cac	cggaggc	cccg	ctttatg	ggtg
151	gaaaacc	ttc	tcatgg	attg	gaagata	ttc	ctgatgt	aga	ggaatat	gag
201	aggaacc	tgc	tcgggg	ctgg	agcaggt	ttt	aatctg	ctca	atgtagg	aaa
251	tatggct	aat	gttccc	gacg	agcacac	acc	gatgat	gtca	ccagtga	ata
301	caactac	aaa	gattct	tacaa	cggagt	ggta	ttaaa	atgga	aatccc	gcca
351	tatttgg	atc	cagacag	tca	ggatgat	gac	ccgga	agatg	gtgtca	acta
401	cccggat	cca	gatttat	tttg	acacaaa	aaa	cacaa	atatg	accgagt	acg
451	atttgg	atgt	gttga	agctt	ggaaa	accag	cagtag	atga	agcacg	gaaa
501	aagatcg	aag	ttccc	gacgc	tagtgc	gccg	caaaca	aaaa	ttgtag	aata
551	tttgatg	tat	tataga	acgt	taaaag	aaa	tgaact	cata	caactga	atg
601	cgtatcg	gac	aaaac	gaaat	cgattat	ctgt	tgaact	ttggt	caaaa	acaat
651	attgatc	gag	agttc	gacca	aaaag	cttgc	gagtc	ccctg	tgaaaa	aat
701	gaaggat	aag	aagaat	gatc	tccaga	acct	gattga	tgtg	gttctt	tcaa
751	aaggtac	aaa	atata	ccggt	tgcatt	tacaa	ttcca	aggac	acttga	tggc
801	cggttac	agg	tccac	ggaag	aaaag	gtttc	cctcac	gtag	tctatg	gcaa
851	actgtgg	agg	ttta	atgaaa	tgacaaa	aaa	cgaa	acgcg	catgtg	gacc
901	actgca	agca	cgcatt	ttgaa	atgaaa	agt	acatg	gtatg	cgtga	atccc
951	tatcact	acg	aaatt	gtcat	tggaa	ctatg	attgt	tgggc	agagg	gatca
1001	tgacaat	cga	gatat	gccgc	cgccac	atca	acgct	tacc	actcc	aggtc
1051	ggcagg	atcc	agttg	acgat	atgag	tagat	ttata	acc	agctt	ccatt
1101	cgtccg	cctc	cgatg	aacat	gcacac	aagg	cctca	gccta	tgcct	caaca
1151	attgc	cctt	ca	gttgg	cga	aa	cgttt	tgcca	tcctc	tcca
1201	cacata	accc	agggg	tttca	catcc	gtact	ccatt	gctcc	acagac	ccat
1251	tacc	cggt	tga	acatg	aaccc	aatcc	cgaa	aa	atgcc	gcaaa
1301	gccacc	acct	ctcc	atcagg	gatat	ggaat	gaatg	gggcc	agttg	ctctt
1351	cagaaa	acaa	caatc	catc	caccaa	aatc	accat	tata	tgat	attag
1401	catcca	aatc	actat	tccta	cgact	gtgg	ccga	actgt	acggg	tttcc
1451	aactc	cttat	ccgg	attttc	accat	ccttt	caat	cagca	ccac	accag
1501	cgccac	aact	atcac	aaaac	cata	cgtccc	aa	caagg	cag	tcat
1551	gggcac	caag	gtcag	gtacc	gaatg	atcca	cca	attt	caa	gacc
1601	acaacc	atca	acagt	cacct	tggac	gtgtt	ccgt	cgtac	tgtag	acaga
1651	cattt	ggaaa	tcg	attttt	gaagg	agaaa	gtga	aca	atc	cggc
1701	attc	ggtcta	gta	acaa	att	cattg	aa	gaa	tttg	attc
1751	tgtg	acagtt	gttc	gaccgc	ggatg	acaga	cgg	t	agg	tt
1801	tcatg	ccgga	agatg	cacca	tatcat	gaca	ttt	g	ca	ag
1851	ctcac	atcag	aaagt	gtaac	tttct	cagga	gag	ggg	ccag	aa
1901	tttga	acgaa	aaatg	ggg	gaa	caatt	gtg	ta	ctat	gagaa
1951	ttggc	gagaa	aaa	atgtt	cg	agagg	aa	att	tcc	acgtgga
2001	attt	gctctg	aga	atcgt	ta	cagtc	ctc	gga	cttg	agccaa
2051	aga	accag	tg	gcgtt	ttaa	ag	ttc	gtaa	agc	aatag
										tggt
										gga
										attc
										gct

Fig. 11B-1



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2101	tttcctacaa	aaaagacggg	agtgtttggc	ttcaaaaccg	catgaagtac
2151	ccggtatattg	tcacttctgg	gtatctcgac	gagcaatcag	gaggcctaaa
2201	gaaggataaa	gtgcacaaag	tttacggatg	tgcgtctatc	aaaacgtttg
2251	gcttcaacgt	ttccaaacaa	atcatcagag	acgcgcttct	ttccaagcaa
2301	atggcaacaa	tgtacttgca	aggaaaattg	actccgatga	attatatcta
2351	cgagaagaag	actcaggaag	agctgcgaag	ggaagcaaca	cgcaccactg
2401	attcattggc	caagtactgt	tgtgtccgtg	tctcgttctg	caaaggattt
2451	ggagaagcat	acccagaacg	cccgtaatt	catgattgtc	cagtttggat
2501	tgagttgaaa	atcaacattg	cctacgattt	catggattca	atctgccagt
2551	acataaccaa	ctgcttcgag	ccgctaggaa	tggaagattt	tgcaaaattg
2601	ggaatcaacg	tcagtgatga	ctaaatgata	acttttttca	ctcaccctac
2651	tagatactga	tttagtctta	ttccaaatca	tccaacgata	tcaaactttt
2701	tcctttgaac	tttgcatact	atggtatcac	aagttccaag	cagtttcaat
2751	acaaacatag	gatatgttaa	caacttttga	taagaatcaa	gttaccaact
2801	gttcattgtg	agctttgagc	tgtatagaag	gacaatgtat	cccatacctc
2851	aatctttaat	agtcatcagt	cactggtccc	gcaccaattt	tttcgattcg
2901	catatgtcat	atattgcacc	gtggcccttt	ttattgtaac	ttttaatata
2951	ttttcttccc	aacttgtgaa	tatgattgat	gaaccaccat	tttgagtaat
3001	aaatgtattt	tttgtgg			

Fig. 11B-2



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1	gtaatcaa	at	tgtaaagg	aa	aatatta	at	agtcagag	ta	cacataa	atg
51	ggtgatc	atc	ataattt	aac	gggcctt	ccc	ggtacct	cca	tcccgcc	aca
101	gttcaac	tat	tctcagc	cccg	gtaccag	cac	cggaggc	ccc	ctttatg	gtg
151	gaaaacc	cttc	tcatgg	attg	gaagata	ttc	ctgatgt	aga	ggaatat	gag
201	aggaacc	tgc	tcgggg	ctgg	agcagg	tttt	aatctgt	ctca	atgtagg	aaa
251	tatggct	aat	gaatttt	aaac	caataat	cac	attggac	acg	aaaccac	ctc
301	gtgatgc	caa	caagtc	attg	gcattca	atg	gcgggtt	gaa	gctaata	cact
351	ccgaaa	actg	aagttcc	cga	cgagcac	aca	ccgatga	tgt	caccagt	gaa
401	tacaact	taca	aagatt	tctac	aacggag	tgg	tattaaa	atg	gaaatcc	cgc
451	catattt	tgga	tccagac	agt	caggatg	atg	acccgga	aga	tggtgtc	aac
501	taccggg	atc	cagattt	tatt	tgacaca	aaa	aacaca	aat	tgaccga	ta
551	cgatttt	ggat	gtggtt	gaagc	ttggaaa	acc	agcagta	gat	gaagcac	gga
601	aaaagat	cga	agttccc	gac	gctagt	gcgc	cgccaa	acaa	aattgt	agaa
651	tatttg	atgt	attatag	aac	gttaaa	agaa	agtga	actca	tacaact	gaa
701	tgcgta	tcgg	acaaa	acgaa	atcgat	tatc	gttga	acttg	gtcaaaa	aca
751	atattg	atcg	agagtt	cgc	caaaa	agctt	gcgag	tcct	ggtgaaa	aaa
801	ttgaag	gata	agaaga	atga	tctcc	agaac	ctgatt	gatg	tggtt	ctttc
851	aaaagg	taca	aaata	taccg	gttgc	attac	aattcc	aagg	acactt	gatg
901	gccgg	ttaca	ggtcc	acgga	agaaa	aggtt	tcctc	acgt	agtct	atggc
951	aaactg	tggga	ggttta	aatga	aatga	caaaa	aacga	aacgc	gtcat	gtgga
1001	ccactg	caag	cacgc	atttg	aatga	aaaag	tgaca	tggta	tgcg	tgaatc
1051	cctatc	acta	cgaa	attgtc	attgg	aaacta	tgatt	gttg	gcag	agggat
1101	catgac	aatc	gagata	tgcc	gccgc	ccacat	caacg	ctacc	acact	ccagg
1151	tcggc	caggat	ccag	ttgacg	atatg	agtag	attta	tacca	ccag	cttcca
1201	ttcgt	ccgcc	tccga	tgaac	atgc	acacaa	ggc	ctcag	cc	tgcctcaa
1251	caattg	cctt	cagtt	ggcgc	aacgt	tttgcc	catc	ctctc	cc	atcaggc
1301	gccaca	ataac	ccagg	gggtt	cacat	ccgta	ctcc	attgct	cc	acagacc
1351	attacc	cggtt	gaac	atgaac	cca	attccg	aaatg	ccgca	aatg	ccacaa
1401	atgcc	accac	ctct	ccatca	ggg	atatgga	atga	atggg	cgag	ttgctc
1451	ttcaga	aaaac	aaca	atccat	tcc	acccaaa	tcacc	attat	aatg	atatta
1501	gccat	ccaaa	tcact	tattcc	tacg	actgtg	gtccg	aactt	gtac	gggttt
1551	ccaact	cctt	atccg	gattt	tcacc	atcct	ttca	atcag	aacc	acacca
1601	gccgc	ccacaa	ctat	cacaaa	acc	atacgtc	cca	acaagg	agtc	atcaac
1651	caggg	cacca	aggt	caggta	ccga	atgatc	cacca	atttc	aag	accagt
1701	ttaca	accat	caac	agtcac	ctt	ggacgtg	ttcc	gtcgg	actg	tagaca
1751	gacatt	tggga	aatc	gatttt	ttga	aggaga	aagt	gaaca	tccg	gcgcaa
1801	taatt	cggtc	tagta	acaaa	ttc	attgaag	aatt	tgat	gcc	gattt
1851	ggtgt	gacag	ttgt	tcgacc	gcgg	atgaca	gacg	gtgag	tttt	ggagaa
1901	catcat	gccg	gaag	atgcac	catat	catga	catt	tgcag	ttc	attttga
1951	ggctc	acatc	agaa	agtgt	acttt	ctcag	gag	agggg	cc	agaagt
2001	gattt	gaacg	aaaa	atggg	aaca	attgtg	tact	atgaga	aaa	atttgca
2051	aattg	gcgag	aaaa	atggt	cgag	aggaaa	tttcc	acgtg	gatg	gcggat

Fig. 11C-1



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2101	tcatttgctc	tgagaatcgt	tacagtctcg	gacttgagcc	aatccaatt
2151	agagaaccag	tggcgtttaa	agttcgtaaa	gcaatagtgg	atggaattcg
2201	cttttcctac	aaaaaagacg	ggagtgtttg	gcttcaaaac	cgcatgaagt
2251	acccggtatt	tgtcacttct	gggtatctcg	acgagcaatc	aggaggccta
2301	aagaaggata	aagtgcacaa	agtttacgga	tgtgcgtcta	tcaaaacggt
2351	tggcttcaac	gtttccaaac	aaatcatcag	agacgcgctt	ctttccaagc
2401	aatgggaac	aatgtacttg	caaggaaaat	tgactccgat	gaattatata
2451	tacgagaaga	agactcagga	agagctgcga	agggaagcaa	cacgcaccac
2501	tgattcattg	gccaagtact	gttgtgtccg	tgtctcgttc	tgcaaaggat
2551	ttggagaagc	ataccagaa	cgcccgctcaa	ttcatgattg	tccagtttgg
2601	attgagttga	aaatcaacat	tgcttacgat	ttcatggatt	caatctgcca
2651	gtacataacc	aactgcttcg	agccgctagg	aatggaagat	tttgcaaaat
2701	tgggaatcaa	cgtcagtgat	gactaaatga	taactttttt	cactcaccct
2751	actagatact	gatttagtct	tattccaaat	catccaacga	tatcaaactt
2801	tttcctttga	actttgcata	ctatgttatc	acaagttcca	agcagtttca
2851	atacaaacat	aggatatgtt	aacaactttt	gataagaatc	aagttaccaa
2901	ctgttcattg	tgagctttga	gctgtataga	aggacaatgt	atcccatacc
2951	tcaatcttta	atagtcatca	gtcactgggc	ccgcaccaat	tttttcgatt
3001	cgcatatgtc	atatattgca	ccgtggccct	ttttattgta	acttttaata
3051	tattttcttc	ccaacttggt	aatatgattg	atgaaccacc	attttgagta
3101	ataaatgtat	tttttgtgg			

Fig. 11C-2



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1	MKLIATSLLV	PDEHTPMMSP	VNTTTKILQR	SGIKMEIPPY	LDPDSQDDDP
51	EDGVNYPDPD	LFDTKNTNMT	EYDLVDLKLK	KPAVDEARKK	IEVPDASAPP
101	NKIVEYLMYY	RTLKESELIQ	LNAYRTKRNR	LSLNLVKNNI	DREFDQKACE
151	SLVKKLKDKK	NDLQNLIDVV	LSKGTKYTGC	ITIPRTL DGR	LQVHGRKGFP
201	HVVYGLWRF	NEMTKNETRH	VDHCKHAFEM	KSDMVCVNPY	HYEIVIGTMI
251	VGQRDHDNRD	MPPPHQRYHT	PGRQDPVDDM	SRFIPPASIR	PPPMNMHTRP
301	QPMPQQLPSV	GATFAHPLPH	QAPHNPGVSH	PYSIAPQTHY	PLNMNPIPQM
351	PQMPQMPPPL	HQGYGMNGPS	CSSENNPFH	QNHHYNDISH	PNHYSYDCGP
401	NLYGFPTYP	DFHHPFNQQP	HQPPQLSQNH	TSQQGSHQPG	HQGOVPNDPP
451	ISRPVLQPST	VTLDVFRRYC	RQTFGNRFFE	GESEQSGAII	RSSNKFIEEF
501	DSPICGVTVV	RPRMTDGEVL	ENIMPEDAPY	HDICKFILRL	TSESVTFSGE
551	GPEVSDLNEK	WGTIVYYEKN	LQIGEKKCSR	GNFHVDGGFI	CSENRYSLGL
601	EPNPIREPVA	FKVRKAIVDG	IRFSYKKDGS	VWLQNRMKYP	VFVTSGYLDE
651	QSGGLKKDKV	HKVYGCASIK	TFGFNVSKQI	IRDALLSKQM	ATMYLQGKLT
701	PMNYIYEKKT	QEELRREATR	TTDSLAKYCC	VRVSFCKGFG	EAYPERPSIH
751	DCPVWIELKI	NIAYDFMDSI	CQYITNCFEP	LGMEDFAKLG	INVSDD

Fig. 12A



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1	MGDHHNLTGL	PGTSIPPQFN	YSQPGTSTGG	PLYGGKPSHG	LEDIPDVEEY
51	ERNLLGAGAG	FNLLNVGNMA	NVPDEHTPMM	SPVNTTTKIL	QRSGIKMEIP
101	PYLDPDSQDD	DPEDGVNYPD	PDLFDTKNTN	MTEYDLDLVK	LGKPAVDEAR
151	KKIEVPDASA	PPNKIVEYLM	YYRTLKESEL	IQLNAYRTRK	NRLSLNLVKN
201	NIDREFDQKA	CESLVKKLKD	KKNDLQNLID	VVLSKGTKYT	GCITIPRTLD
251	GRLQVHGRKG	FPHVVYGLW	RFNEMTKNET	RHVDHCKHAF	EMKSDMVCVN
301	PYHYEIVIGT	MIVGQRDHDN	RDMPPPHQRY	HTPGRQDPVD	DMSRFIPPAS
351	IRPPPMNMHT	RPQPMPOQLP	SVGATFAHPL	PHQAPHNPGV	SHPYSIAPQT
401	HYPLNMNPIP	QMPQMPQMP	PLHQGYGMNG	PSCSSENNNP	FHQNHHYNDI
451	SHPNHYSYDC	GPPLYGFPTP	YPDFHHPFNQ	QPHQPPQLSQ	NHTSQQGS HQ
501	PGHQGQVPND	PPISRPVLQP	STVTLDVFRR	YCRQTFGNRF	FEGESEQSGA
551	IIRSSNKFIE	EFDSPICGVT	VVRPRMTDGE	VLENIMPEDA	PYHDICKFIL
601	RLTSESVTFS	GEGPEVSDLN	EKWGTIVYYE	KNLQIGEKKC	SRGNFHV DGG
651	FICSENRYSL	GLEPNPIREP	VAFKVRKAIV	DGIRFSYKKD	GSVWLQNRMK
701	YPVFVTSGYL	DEQSGGLKKD	KVHKVYGCAS	IKTFGFNVSK	QIIRDALLSK
751	QMATMYLQ GK	LTPMNYIYEK	KTQEELRREA	TRTTDSLAKY	CCVRVSFCKG
801	FGEAYPERPS	IHDCPVWIEL	KINIAYDFMD	SICQYITNCF	EPLGMEDFAK
851	LGINVSDD				

Fig. 12B



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1	MGDHHNLTGL	PGTSIPPQFN	YSQPGTSTGG	PLYGGKPSHG	LEDIPDVEEY
51	ERNLLGAGAG	FNLLNVGNMA	NEFKPIITLD	TKPPRDANKS	LAFNGGLKLI
101	TPKTEVPDEH	TPMMSPVNTT	TKILQRSGIK	MEIPPYLDPD	SQDDDPEDGV
151	NYPDPDLFDT	KNTNMTEYDL	DVLKLGKPAV	DEARKKIEVP	DASAPPNKIV
201	EYLMYYRTLK	ESELIQLNAY	RTKRNRSLN	LVKNNIDREF	DQKACESLVK
251	KLKDKKNDLQ	NLIDVVLSKG	TKYTGCTIP	RTLDGRLQVH	GRKGFPHVY
301	GKLWRFNEMT	KNETRHVDHC	KHAFEMKSDM	VCVNPYHYEI	VIGTMIVGQR
351	DHDNRDMPPP	HQRYHTPGRQ	DPVDDMSRFI	PPASIRPPPM	NMHTRPQMP
401	QQLPSVGATF	AHPLPHQAPH	NPGVSHPYSI	APQTHYPLNM	NPIPQMPQMP
451	QMPPPLHQGY	GMNGPSCSSE	NNNPFHQNH	YNDISHPNHY	SYDCGPNLYG
501	FPTPYPDFHH	PFNQPHQPP	QLSQNHTSQQ	GSHQPGHQGQ	VPNDPPISRP
551	VLQPSTVTLD	VFRRYCRQTF	GNRFFEGESE	QSGAIIRSSN	KFIEEFDSP
601	CGVTVVRPRM	TDGEVLENIM	PEDAPYHDIC	KFILRLTSES	VTFSGEGPEV
651	SDLNEKWGTI	VYYEKNLQIG	EKKCSRGNFH	VDGGFICSEN	RYSLGLEPNP
701	IREPVAFKVR	KAIVDGIRFS	YKKGDSVWLQ	NRMKYPVFT	SGYLDEQSGG
751	LKKDKVHKVY	GCASIKTFGF	NVSKQIIRDA	LLSKQMATMY	LQKLTMPNY
801	IYEKKTQEEL	RREATRTTDS	LAKYCCVRVS	FCKGFGEAYP	ERPSIHDCPV
851	WIELKINIAY	DFMDSICQYI	TNCFEPLGME	DFAKLGINVS	DD

Fig. 12C

Fig. 13A

ttacacgtggccaatgcaacaatacatctatcaggaatcgtcagcaaccattccccatcaccattttaaatcaacacaaca
atccgtatcatccaatgcatcctcatcatcaattacctcatatgcaacaacttcccaacctctattgaaatcttaacatg
acgacgttaacatcttctggcagttccgtggccagttccattggaggcggagctcaatgctctccgtgcgcgtcgggctc
ctcgaccgtgcaacaaattcctctcaacagcagcagaccgttggtcaaatgcttgctgcacgtgccttgttcttcat
ctggcatgacacttggaatgtcacttaatctgtcacaaggcgggtggccaatgccggcaaaaaagaagcgttgctcgaag
aagccaaccgatcaattggcacagaagaaaccgaatccatggggtgaggaatcctattcgggatcattgccaagcatt
ggaatcggcgccagacggaaggttaaactcaatgagatttatcaatgggttctctgataatattccctactttggagaac
gatctagtcgggaggggccgcccggatggaagaactcgatccgtcacaatctgtctcttctcattctcgtttcatgcaatt
cagaatgaaggagccggaagagctcgtgggtgggttattaatccagatgcaaagccaggaatgaatccacggcgtacacg
tgaacgatccaatactattgagacgactacaaaggctcaactcgaaaaatctcgccgggagccaagaagaggataaagg
agagagcattgatgggctcccttctcactcgacacttaatggaaatcgattgccggatcgattcaaacgatttctcacgat
ttgtatgatgatgattcaatgcaaggagcatttgataacgttccatcatcttccgtccccgaactcaatcgaaacctctc
gattcctggatcgtcgtctcgtgttctccagctattggaagtgatattctatgatgatctagaattcccatcatgggttg
gcaatcgggtccagcaattccaagtgatattgttgatagaactgatcaaatgcgtatcgatgcaactactcatattgggt
ggagttcagattaagcaggagtcgaagccgattaaagacggaaccaattgctccaccaccatcataccacgagttgaacag
tgtccgtggatcgtgtgctcagaatccacttcttcgaaatccaattgtgccaagcactaacttcaagccaatgccactac
cgggtgcctatggaaactatcaaaatgggtggaataactccaatcaattggctatcaacatccaactcatctccactgcct
ggaattcaatcgtgtggaattgtagctgcacagcatactgtcgtcttctcatcggtcttccaattgatttggaataatct
gacacttcccgatcagccactgatggatactatggatgttgatgcattgatcagacatgagctgagtcgaagctggagggc
agcatattcattttgatttgtaaattctcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttctt
ggagtgagaaatcttccgtcttctcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttctt
cttctctctcgtctaatccaacacattcatcccagtgacgtcgtgtaataataataataataacacctctctctcttctt
ccctaatgcgaaatatcgaaaaaccgttgattattacctcttcttcttcttcttcttcttcttcttcttcttcttctt
tccaggttcttctactctttaaattgctacctctatcccatcttcttcttcttcttcttcttcttcttcttcttcttct
acacattccccaatctgtctt
tctctt
gaatcctcgtatacacacacacatagtaattctacctccaaaattttactgaaagatgtgatccccctctctgtctccctc
tacaaaacattatttctgtgttctgtgtatattgccaccacgtcgatttttaattaaaaccatcggttttcttcttcttct
acttttctctcgaaaaatttaacaacacacaaaaaaatccttcaaaaaatctcagttttaaatgggtgtggcaatatatcg
gatccccctctacaccagaacagtccttgcaatttcagagaaatgattttcagatttttcatatcacaggccccctttttt
gcttggtttttctctacctctcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttcttctt
ttccaattctt
tcgtctccctccgcccccaatatatttgcgactgtatgatgatgatgatgatttaataaaaaat

Fig. 13B



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MMEMLVDQGTDASSSASTSTSSVSFRGADTFMNTPDVVMMNDDMEPIPRDR
CNTWPMRRPQLEPPLNSSPIIHEQIPEEDADLYGSNEQCGQLGGASSNGST
AMLHTPDGSNSHQTSFSPSDFRMSSEPD DTVSGKKTTTRRNAWGNMSYAEI
TTAIMASPEKRLTLAQVYEWVQNVFYFRDKGDSNSSAGWKNSIRHNLSLH
SRFMRIQNEGAGKSSWWVINPDAKPGMNPRRTRERSNTIETTTKAQLEKSR
RGAKKRIKERALMGSLHSTLNGNSIAGSIQTISHDLYDDDSMQGAFDNVPS
SFRPRTQSNLSIPGSSSRVSPAIGSDIYDDLEFPSWVGESVPAIPSDIVDR
TDQMRIDATTHIGGVQIKQESKPIKTEPIAPPPSYHELNSVRGSCAQNP LL
RNPIVPSTNFKPMPLPGAYGNYQNGGITPINWLSTSNSSPLPGIQSCGIVA
AQHTVASSALPIDLENLTLPDQPLMDTMDVDALIRHEL SQAGGQHIHFDL

Fig. 14A

MQQYIYQESSATIPHHHLNQHNPNYPHMPHHQLPHMQQLPQPLLNLNMTT
LTSSGSSVASSIGGGAQCSPCASGSSTAATNSSQQQQQTVGQMLAASVPCSS
SGMTLGMSLNLSQGGGPMPAKKKRCKKPTDQLAQKKPNPWGEESYSDIIA
KALESAPDGRKLKLEIYQWFSNIPYFGERSSP EEAAGWKNSIRHNLSLHS
RFMRIQNEGAGKSSWWVINPDAKPGMNPRRTRERSNTIETTTKAQLEKSRR
GAKKRIKERALMGSLHSTLNGNSIAGSIQTISHDLYDDDSMQGAFDNVPS
FRPRTQSNLSIPGSSSRVSPAIGSDIYDDLEFPSWVGESVPAIPSDIVDR
DQMRIDATTHIGGVQIKQESKPIKTEPIAPPPSYHELNSVRGSCAQNP LLR
NPIVPSTNFKPMPLPGAYGNYQNGGITPINWLSTSNSSPLPGIQSCGIVAA
QHTVASSALPIDLENLTLPDQPLMDTMDVDALIRHEL SQAGGQHIHFDL

Fig. 14B



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1	cggaagccat	ggagctcgag	atctgattgc	tggacacgga	cggaactccg	acgtatctcg
61	cagatgcatg	ttaacatttt	acatccacaa	ctgcaaacga	tggctcgagca	gtggcaaagt
121	cgagaacgcc	catcgctgga	gaccgagaat	ggcaaaggat	cgctgctcct	ggaaaatgaa
181	ggtgtcgag	atatcatcac	tatgtgtcca	ttcggagaag	ttattagtgt	agtatttccg
241	tggtttcttg	caaatgtgcy	aacatcgcta	gaaatcaagc	tatcagattt	caaacatcaa
301	cttttcgaat	tgattgctcc	gatgaagtgg	ggaacatatt	ccgtaaagcc	acaggattat
361	gtgttcagac	agttgaataa	tttcggcgaa	attgaagtta	tatttaacga	cgatcaaccc
421	ctgtcgaaat	tagagctcca	cggcactttc	ccaatgcttt	ttctctacca	acctgatgga
481	ataaacaggg	ataaagaatt	aatgagtgat	ataagtcatt	gtctaggata	ctcactggat
541	aaactggaag	agagcctcga	tgaggaactc	cgtcaatttc	gtgcttctct	ctgggctcgt
601	acgaagaaaa	cgtgcttgac	acgtggactt	gagggtagca	gtcactacgc	gttccccgaa
661	gaacagtact	tgtgtgttgg	tgaatcgctc	ccgaaagatt	tggaaatcaaa	agtcaaggct
721	gccaagctga	gttatcagat	gttttgagga	aaacgtaaa	cggaatcaaa	tggagtttgc
781	gagaaaatga	tgaagattca	aattgaattc	aatccgaacg	aaactccgaa	atctctgctt
841	cacacgtttc	tctacgaaat	gcgaaaattg	gatgtatacg	ataccgatga	tcctgcagat
901	gaaggatggg	ttcttcaatt	ggctggacgt	accacgtttg	ttacaaatcc	agatgtcaaa
961	cttacgtctt	atgatgggtg	ccgttcggaa	ctggaaagct	atcgatgccc	tggattcgtt
1021	gttcgcccga	aatcactagt	cctcaaagac	tattgtcgcg	caaaaccact	ctacgaacca
1081	cattatgtga	gagcacacga	acgaaaactt	gctctagacg	tgctcagcgt	gtctatagat
1141	agcacaccaa	aacagagcaa	gaacagtgcg	atgggttatga	ctgatttttcg	tccgacagct
1201	tactcaaac	aagtttcact	ttgggacctt	gacgcgaatc	ttatgatacg	gcctgtgaat
1261	atttctggat	tcgatttccc	ggccgacgtg	gatatgtacg	ttcgaatcga	attcagtgtg
1321	tatgtgggga	cactgacgct	ggcatcaaaa	tctacaacaa	aagtgaatgc	tcaatttgca
1381	aaatggaata	aggaaatgta	cacttttgat	ctatacatga	aggatatgcc	accatctgca
1441	gtactcagca	ttcgtgtttt	gtacggaaaa	gtgaaattaa	aaagtgaaga	attcgaagtt
1501	ggttgggtaa	atatgtccct	aaccgattgg	agagatgaac	tacgacaagg	acaattttta
1561	ttccatctgt	gggctcctga	accgactgcc	aatcgtagta	ggatcggaga	aaatggagca
1621	aggataggca	ccaacgcagc	ggttacaatt	gaaatctcaa	gttatgggtg	tagagttcga
1681	atgccgagtc	aaggacaata	cacatatctc	gtcaagcacc	gaagtacttg	gacggaaact
1741	ttgaatatta	tgggtgatga	ctatgagtcg	tgtatcagag	atccaggata	taagaagctt
1801	cagatgcttg	tcaagaagca	tgaatctgga	attgtattag	aggaagatga	acaacgtcat
1861	gtctggatgt	ggaggagata	cattcaaaa	caggagcctg	atttgctcat	tgtgctctcc
1921	gaactcgcat	ttgtgtggac	tgatcggtgag	aacttttccg	agctctatgt	gatgcttgaa
1981	aaatggaaac	cgccgagtg	ggcagccg	ttgactttgc	ttggaaaacg	ttgcacggat
2041	cgtgtgattc	gaaagtgtgc	agtggagaag	ttgaatgagc	agctgagccc	ggtcacattc
2101	catcttttca	tattgcctct	catacaggcg	ttgaagtacg	aaccgcgtgc	tcaatcgga
2161	gttggaaatga	tgctcttgac	tagagctctc	tgcgattatc	gaattggaca	tcgacttttc
2221	tggctgctcc	gtgcagagat	tgctcgtttg	agagattgtg	atctgaaaag	tgaagaatat
2281	cgccgtatct	cacttctgat	ggaagcttac	ctccgtggaa	atgaagagca	catcaagatc
2341	atcacccgac	aagttgacat	ggttgatgag	ctcacacgaa	tcagcactct	tgtcaaagga
2401	atgccaaaag	atgttgctac	gatgaaactg	cgtgacgagc	ttcgatcgat	tagtcataaa
2461	atggaaaata	tggattctcc	actggatcct	gtgtacaaac	tgggtgaaat	gataatcgac
2521	aaagccatcg	tcctaggaag	tgcaaaacgt	ccgttaatgc	ttcactggaa	gaacaaaaat
2581	ccaaagagtg	acctgcacct	tccgttctgt	gcaatgatct	tcaagaatgg	agacgatctt
2641	cgccaggaca	tgcttgttct	tcaagttctc	gaagttatgg	ataacatctg	gaaggctgca

Fig. 15-1



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2701	aacattgatt	gctgtttgaa	cccgtagca	gttcttccaa	tgggagaaat	gattggaatt
2761	attgaagttg	tgcctaattg	taaaacaata	ttcgagattc	aagttggaac	aggattcatg
2821	aatacagcag	ttcggagtat	tgatccttcg	tttatgaata	agtggattcg	gaaacaatgc
2881	ggaattgaag	atgaaaagaa	gaaaagcaaa	aaggactcta	cgaaaaatcc	catcgaaaag
2941	aagattgata	atactcaagc	catgaagaaa	tattttgaaa	gtgtcgatcg	attcctatac
3001	tcgtgtgttg	gatattcagt	tgccacgtac	ataatgggaa	tcaaggatcg	tcacagtgat
3061	aatctgatgc	tcactgaaga	tggaaaatat	gtccacattg	atttcggtca	cattttggga
3121	cacggaaaga	ccaaacttgg	gatccagcga	gatcgtcaac	cgtttattct	aaccgaacac
3181	tttatgacag	tgattcgatc	gggtaaatct	gtggatggaa	attcgcata	gctacaaaaa
3241	ttcaaaacgt	tatgcgtcga	agcctacgaa	gtaatgtgga	ataatcgaga	tttgttcgtt
3301	tccttgttca	ccttgatgct	cggaatggag	ttgcctgagc	tgtcgacgaa	agcggatttg
3361	gatcatttga	agaaaaccct	cttctgcaat	ggagaaagca	aagaagaagc	gagaaagttt
3421	ttcgctggaa	tctacgaaga	agccttcaat	ggatcatggt	ctacccaaaac	gaattggctc
3481	ttccacgcag	tcaaactacta	ctga			

Fig. 15-2



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1	RKPWSSRSDC	WTRTELRRIS	QMHVNILHPQ	LQTMVEQWQM	RERPSLETEN	GKGSLLLENE
61	GVADIITMCP	FGEVISVVP	WFLANVRTSL	EIKLSDFKHQ	LFELIAPMKW	GTYSVKPODY
121	VFRQLNNFGE	IEVIFNDDQP	LSKLELHGTF	PMLFLYQPDG	INRDKELMSD	ISHCLGYSLD
181	KLEESLDEEL	RQFRASLWAR	TKKTCLTRGL	EGTSHYAFPE	EQYLCVGESC	PKDLESKVKA
241	AKLSYQMFWR	KRKAEINGVC	EKMMKIQIEF	NPNETPKSLL	HTFLYEMRKL	DVYDTDDPAD
301	EGWFLQLAGR	TTFVTNPDVK	LTSYDGVRS	LESYRCPGFV	VRRQSLVLKD	YCRPKPLYEP
361	HYVRAHERKL	ALDVLSVSID	STPKQSKNSD	MVMTDFRPTA	SLKQVSLWDL	DANLMIRPVN
421	ISGFDFPADV	DMYVRIEFSV	YVGTTLTASL	STTKVNAQFA	KWNKEMYTFD	LYMKDMPPSA
481	VLSIRVLYGK	VKLKSEEFV	GWVNMSLTDW	RDELROGQFL	FHLWAPEPTA	NRSRIGENGA
541	RIGTNAAVTI	EISSYGGVR	MPSQGQYTYL	VKHRSTWTET	LNIMGDDYES	CIRDPGYKKL
601	QMLVKKHESG	IVLEEDEQRH	VWMWRRYIQK	QEPDLLIVLS	ELAFVWTDRE	NFSELYVMLE
661	KWKPPSVAAA	LTLLGKRCTD	RVIRKFAVEK	LNEQLSPVTF	HLFILPLIQA	LKYEPRQSE
721	VGMMLLTRAL	CDYRIGHRLF	WLLRAEIARL	RDCDLKSEY	RRISLLMEAY	LRGNEEHIKI
781	ITRQVDMVDE	LTRISTLVKG	MPKDVATMKL	RDELRSISHK	MENMDSPLDP	VYKLGEMIID
841	KAIVLGSARK	PLMLHWKNKN	PKSDLHLPFC	AMIFKNGDDL	RQDMLVLQVL	EVMDNIWKAA
901	NIDCCLNPYA	VLPNGEMIGI	IEVVPNCKTI	FEIQVGTGFM	NTAVRSIDPS	FMNKWIRKQC
961	GIEDEKKKSK	KDSTKNPIEK	KIDNTQAMKK	YFESVDRFLY	SCVGYSVATY	IMGIKDRHSD
1021	NLMLTEDGKY	VHIDFGHILG	HGKTKLGIQR	DRQPFILTEH	FMTVIRSGKS	VDGNSHELQK
1081	FKTLCVEAYE	VMWNNRDLFV	SLFTLMLGME	LPELSTKADL	DHLKKTLCFN	GESKEEARKF
1141	FAGIYEEAFN	GSWSTKTNL	FHAVKHY			

Fig. 16

**CONVERGENT TGF- β AND INSULIN SIGNALING
ACTIVATE GLUCOSE-BASED METABOLISM GENES**

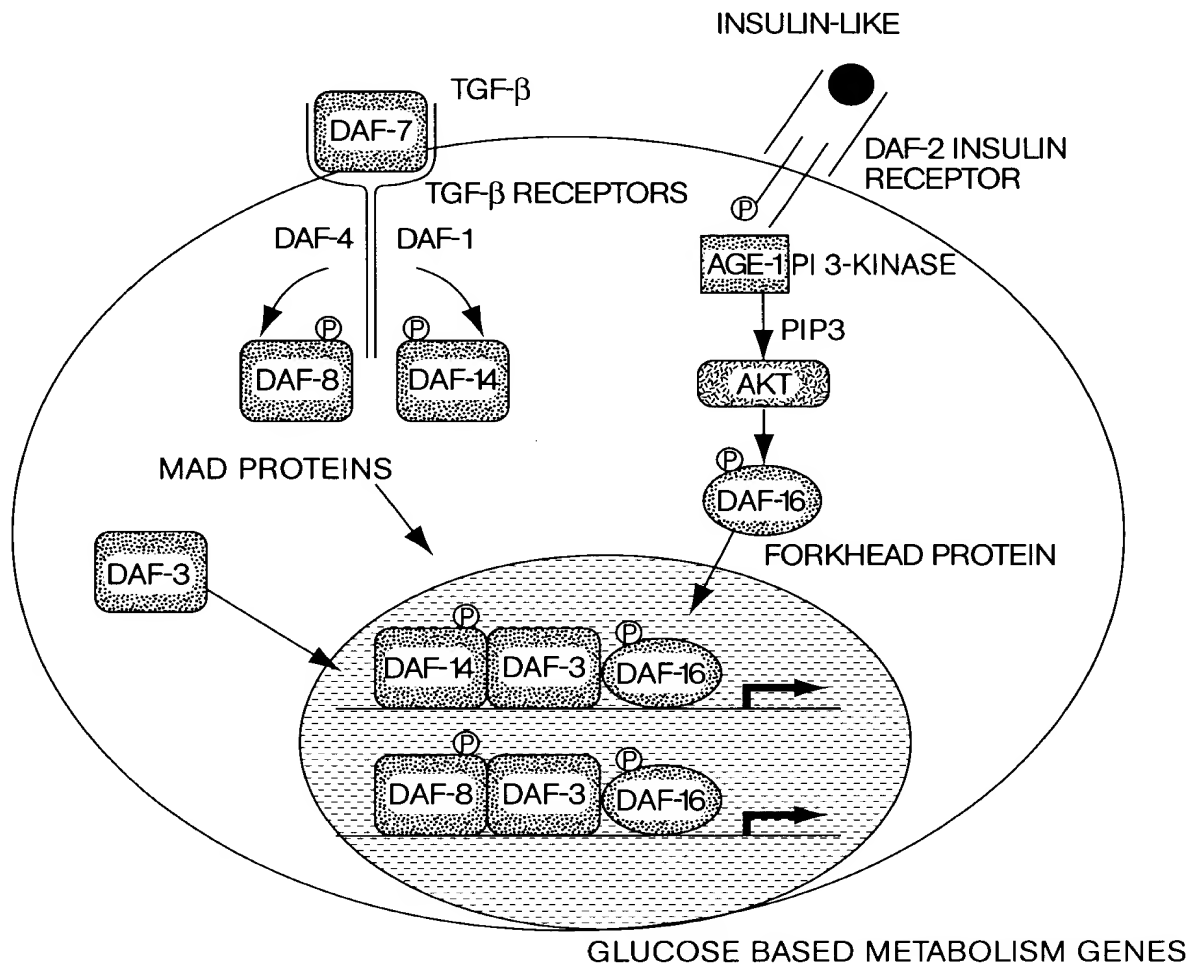


Fig. 17

**IN PHEROMONE, NO TGF β OR INSULIN-LIKE SIGNALS
CAUSES REPRESSION OF ANABOLIC GENES**

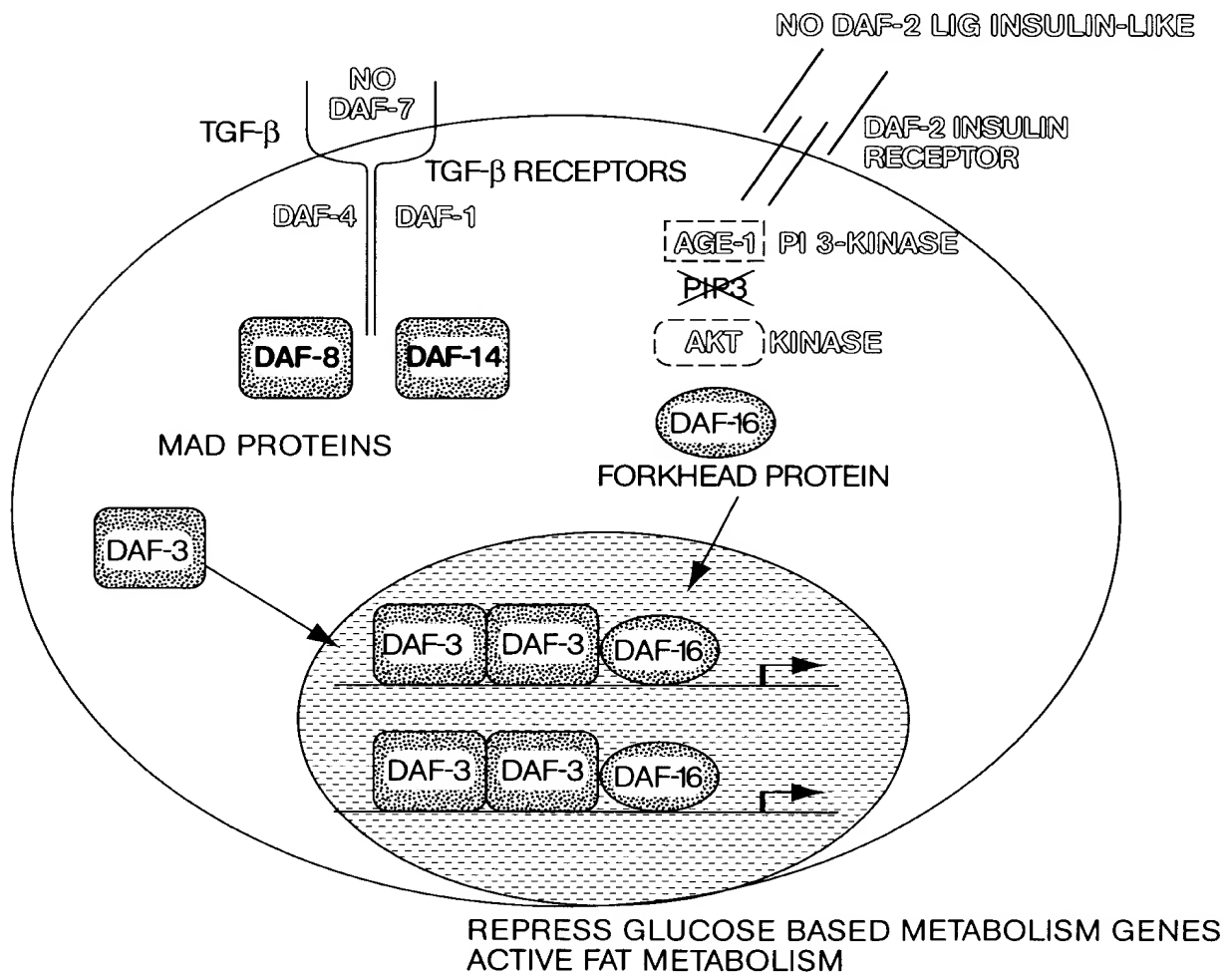
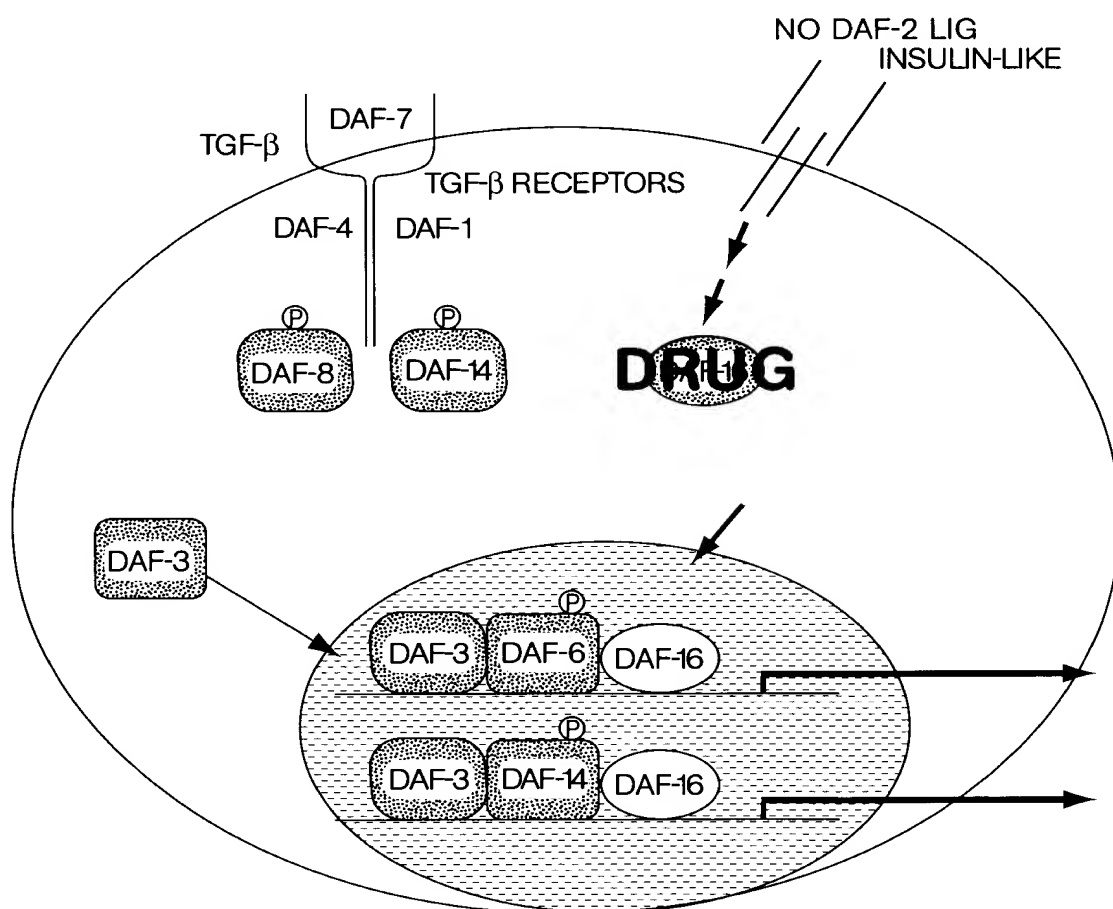


Fig. 18

**DRUGS THAT INHIBIT DAF-16 OR DAF-3
(OR PROTEINS IN THE PATHWAY)
CAN BE DISCOVERED USING REPORTER GENES
BEARING THEIR COGNATE BINDING SITES**



DRUG CAUSES A DECREASE IN DAF-16 ACTIVITY, ACTIVATING
THE REPORTER GENE LIKE A DAF-16 MUTANT.
THIS BYPASSES THE NEED FOR INSULIN

Fig. 19

**DRUGS THAT INHIBIT DAF-3 WILL CURE
 THE DIABETES CAUSED BY A LACK OF DAF-7**

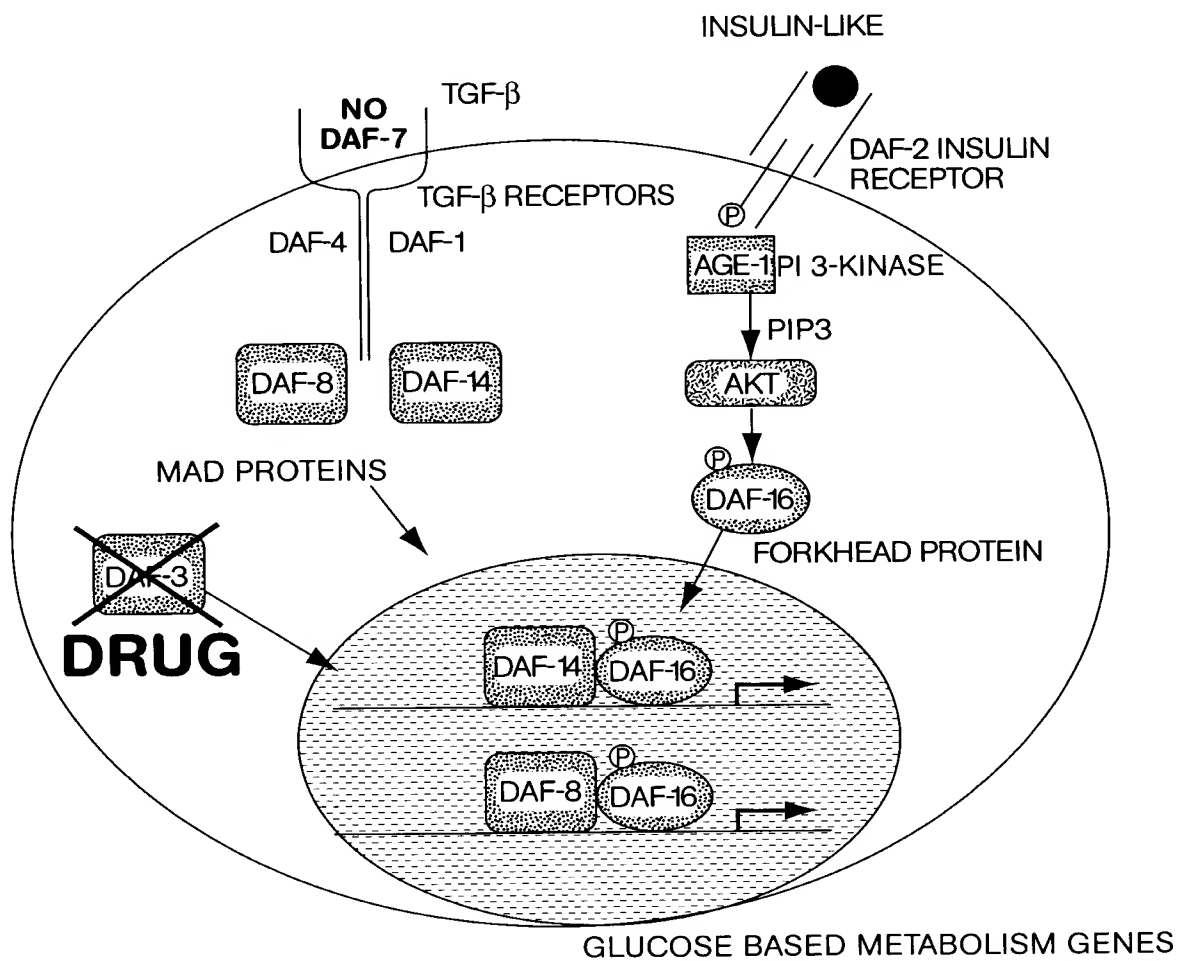


Fig. 20



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DAF-16a1 1 ---MMMLVDQGTDASSSASTSTSSVRFQADTFMNTDDVMNDDEMEIPRDR
DAF-16b 1 ---MNDSTDDPEPEPRGRCTWPMQYIYQESSATIPHHLNQHNNEYPHMPHHQLPHMQLPQELLN
FKHR 1 ---MAEAPQVWEIDDEPEPLPRPRSCWPLPRPEFQSQNSATSPAPSGSAAN.....DAAAGLPSASA
FKHRL1 1 MAEAPAPLSPLEVELDEPEPEPRSCWPLPRPELQASPAKPGETAADSMIPE.....EEDDEDEDGG
AFX 1 ---MRIOBQKAA

DAF-16a1 52 CN..WPMRRRPOLEPPNSPIIHEQIPEEDADLYCSNEQ...CCOLGASNGSTAMHTPDGNSHOTSFFSDFRMSE
DAF-16b 68 LNMTLTSSGSSVASSIGGAQCSPCASGSSTATNSQQQQTQVQOMLAAVPCSSSGMTGMSINLSOGGPMPAKKR
FKHR 64 AAVSADFMNSNLSTLEESDFQAPGSVAANAFAAAAAATGGLGDFQGPAGEG..LHPAPPOPPPGGLSQHPPVPPAA
FKHRL1 72 RAGSAMAIGGGGSGTGGLLEDS..ARVLAPEGQDPCGCPATAAGGLSGGT..QALLQPOOPLP.....PPQPGAAG
AFX 10 AIIDLDDDFEPRSRSCWPLPRPEIANQPSPEPEVEDLGEKVHTEGSEPI..LPSRUSEPAGGE...QPGILCAVT

DAF-16a1 127 SPDDTVSGKKTTRNNAWGNMSVAELTITAITAMASPEKRLTLAQVEMWVQNPVPERDKGDSNSAGWKNSTRHNSLSHSR
DAF-16b 148 CRKKP.TDOLAQKKPNWGEESVSDITAKALETSAPDGRKLNEMVOWFSNDNIPYFGERSSPEEAAGWKNSTRHNSLSHSR
FKHR 143 GPLAGOPRKSSSRNAGNLSVADLITKAIESAEKRLTISOIMEMVKSVPYFKDKGDSNSAGWKNSTRHNSLSHSK
FKHRL1 143 G..SGOPRK.CSSRRNAGNLSVADLITRAIESSPDKRLTISOIMEMVRCVPYFKDKGDSNSAGWKNSTRHNSLSHSR
AFX 86 GPRKC.....GSRNAGNLSVAEFTISOAIESAPEKRLTLAQVEMWVTVPMYFKDKGDSNSAGWKNSTRHNSLSHSK

DAF-16a1 207 FMRIONEGAGKSSWVWVNDPAKPGRNPRRTRERSNTIETITKAOLEKSRGAKRKIRALMGSLHSTNGNSTAGSIOT
DAF-16b 227 FMRIONEGAGKSSWVWVNDPAKPGRNPRRTRERSNTIETITKAOLEKSRGAKRKIRALMGSLHSTNGNSTAGSIOT
FKHR 223 FIRVONEGTGKSSWVWVNDPAKPGRNPRRTRERSNTIETITKAOLEKSRGAKRKIRALMGSLHSTNGNSTAGSIOT
FKHRL1 220 FMRVONEGTGKSSWVWVNDPAKPGRNPRRTRERSNTIETITKAOLEKSRGAKRKIRALMGSLHSTNGNSTAGSIOT
AFX 160 FIKVHNEATGKSSWVWVNDPAKPGRNPRRTRERSNTIETITKAOLEKSRGAKRKIRALMGSLHSTNGNSTAGSIOT

DAF-16a1 287 ISHDL.YDDDSMOGAFDNPSSFRPRRTOSNLS..PGSSRSVSPATGSDIYDDDL..EPPSWVGESVPAIPSDIVDR.TDOMRIDA
DAF-16b 307 ISHDL.YDDDSMOGAFDNPSSFRPRRTOSNLS..PGSSRSVSPATGSDIYDDDL..EPPSWVGESVPAIPSDIVDR.TDOMRIDA
FKHR 292 FSKWPAASPGSHSNDDFDNPSSFRPRRTOSNLS..PGSSRSVSPATGSDIYDDDL..EPPSWVGESVPAIPSDIVDR.TDOMRIDA
FKHRL1 288 LSKWPGSPTSRSSDELDAWTFERSRTSNAS..TISGRISPM..TFODDLCEGD..MHSWVYPPSAAKMASI.....
AFX 231 FAKWSGSPCSRNEEADMTTTERPRSSNAS..SVSTRFSLRPESEV.LAEIIPASVSSYAGGVPTLNEGLELLDGLN

DAF-16a1 366 TTHIGGVQIKOESKPIKTEPIAPPSSVHELNSVRGSCAONPLTRNPTVPSITNEKPMPLPGAGVGNVONGGITPINWLSISN
DAF-16b 386 TTHIGGVQIKOESKPIKTEPIAPPSSVHELNSVRGSCAONPLTRNPTVPSITNEKPMPLPGAGVGNVONGGITPINWLSISN
FKHR 359 TPLSSEISNPNM.ENLLDNL.NLSSPTSLTSTQSSPGTMMQOTPCYSFAPP.NTSLNSPSPNYOKYTYGQSSMSPP
FKHRL1 366 TPLRTDMAGTMNNDGLTENLMDLLDNITLPPSQSPSTGIMQSSSEPYTTK.GSGLGSPSSFNSTVFGPSSLSNR
AFX 308 TSSSHLLSRSGESGFSLQHPGVGTGLTYSSSLFSPAEGPLSAGEGCSSSQALEALTSDPPPPADVLMTOVDPIIS

DAF-16a1 446 SSPLPGIOS..CGIVAAOHTVASSALPIDENLTUPDOPLMDTMDVDALEIRHELSQAGGOHHEFDL
DAF-16b 466 SSPLPGIOS..CGIVAAOHTVASSALPIDENLTUPDOPLMDTMDVDALEIRHELSQAGGOHHEFDL
FKHR 436 OMPLOTLODNK..SSYGMSQYNCAPEGLKELTSDSRPHNDI.MTPVDPGVAQPNRVLGNV...MGPNSVMSTYGSQ
FKHRL1 445 QSPMQTLOENKPAFFSSMSHY..GNQTLQDLE.TSDLSHSDVMTQSDPLMSQASTAVSAQNSRNRVMLRNDPMSFAAQ
AFX 388 OAPTLLLLLGLPSS....SKLATGVGLCPKPEAREGSSSLVPTLSMIAEPVPMASAPIPKALGTPVLTPEEASQDRMP

Fig. 21A-1



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DAF-16a1	511	-----
DAF-16b	531	-----
FKHR	511	ASHNKMNPSH.THPCHAOQTSAVNCRPIPHIVSTMBHTSGMNRITQVKTPVQVPLPHPMQMSALGGYSSVSSCNGYGR
FKHRL1	523	PNQGS1VN.QN1L.LHQHOTQGALGGSRAFSNSVSNM.GLSSESSLGSAHQOQS2VSQSMQ.TLSDSLGSSLYSTSAN
AFX	464	QDLDLDMYMENFECDMDNIIISDLMEGEGEDFNFEPP
DAF-16a1	511	-----
DAF-16b	531	-----
FKHR	590	MGLLHQEKLPSSDLD.GMEIERLDCDMESTIRNDLMBGDTEDFNFDNVLPNQ.....SEPHSVKTTTHSWVSC
FKHRL1	599	LPVMGHEKFPSSDLDLDMENGSLFCDMESIRSEIMDADGLDFNFDLISTQNVVGLNVGNFTGAKQASSQSWVPC
AFX	502	-----

Fig. 21A-2

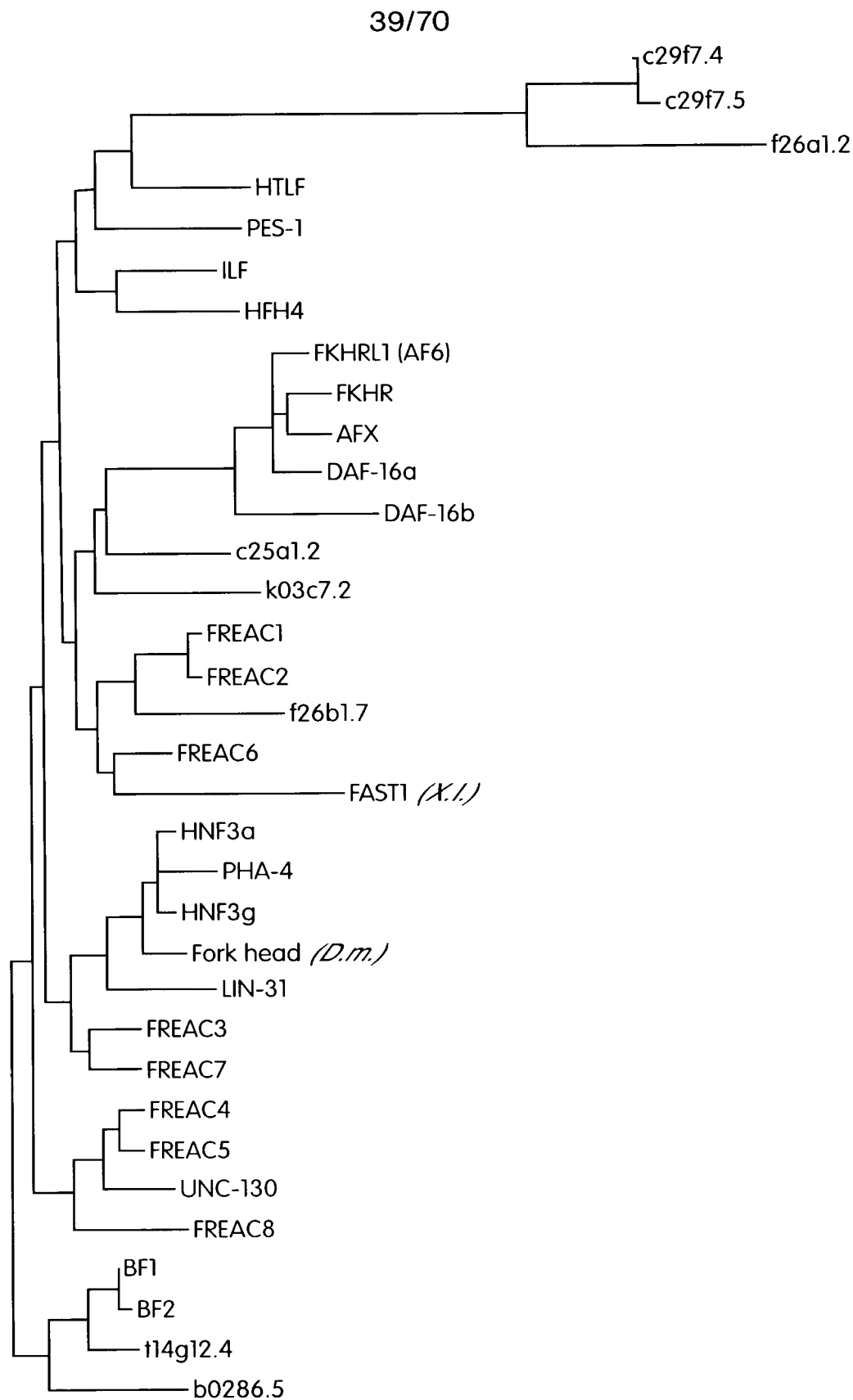


Fig. 21B



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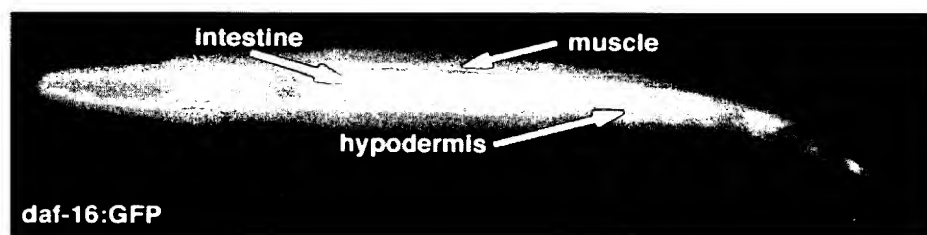


Fig. 22

INJECTION OF OF DAF-7 BYPASSES OBESITY-INDUCED DEFECTS IN INSULIN-REGULATION OF METABOLISM

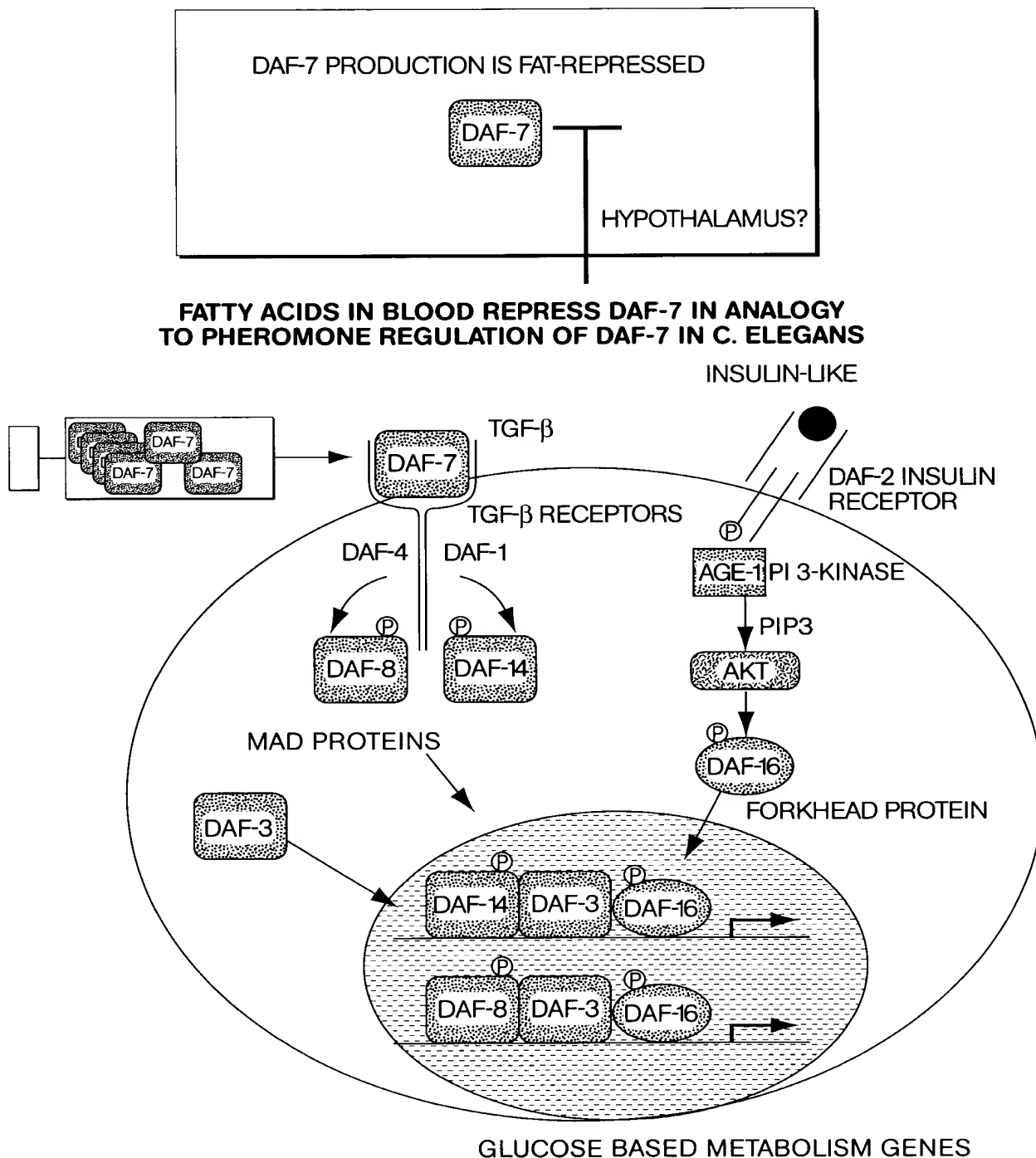


Fig. 23



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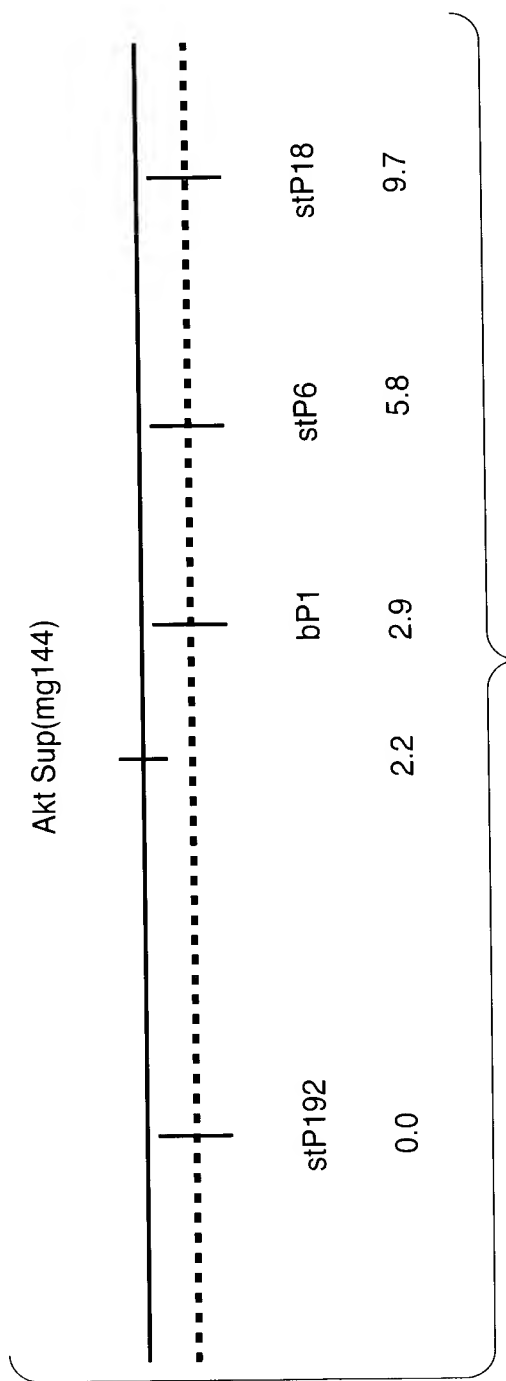
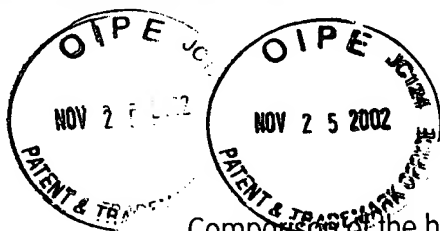


Fig. 24



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Comparison of the human AKT protein sequence to the cosmid sequence C12D8, located in the genetic interval where sup(mgl44) maps. Numbering in the AKT protein sequence by amino acid residues, and in the cosmid sequence by nucleotide position.

Score = 450 (207.4 bits), Expect = 5.2e-165, Sum P(7) = 5.2e-165
Identities = 79/121 (65%), Positives = 97/121 (80%), Frame = +1

Query: 319 EVLEDNDYGRAVDWWGLGVVEMYEMMCGRLPFYNDHEKLFELILMEEIRFPRTLGPPEAKS 378
+VL+D+DYGR VDWVG+GVVEMYEMMCGRLPFY++DH KLFELI+ ++RFP L EA++
Sbjct: 33685 QVLDDHDYGRCDVWVGWGVVEMYEMMCGRLPFYSKDHNKLFELIMAGDLRFPSKLSQEART 33864

Query: 379 LLSGLLKKDPTQRLGGGSEDAKEIMQHRFFANIVWQDVYEKKLSPPFKPQVTSETDTRYFD 439
LL+GLL KDPTQRLGGG EDA EI + FF + W+ Y K++ PP+KP V SETDT YFD
Sbjct: 33865 LLTGLLVKDPTQRLGGGPEDALEICRADFFRTVDWEATYRKEIEPPYKPNVQSETDTSYFD 34047

Score = 256 (118.0 bits), Expect = 5.2e-165, Sum P(7) = 5.2e-165
Identities = 48/66 (72%), Positives = 59/66 (89%), Frame = +1

Query: 146 TMNEFEYLKLLGKGTFGKVILVKEKATGRYYAMKILKKEVIVAKDEVAHTLTENRVLQNS 205
TM +F++LK+LGKGTFGKVIL KEK T + YA+KILKK+VI+A++EVAHTLTENRVLQ
Sbjct: 32314 TMEDFDLKVILGKGTFGKVILCKEKRTQKLYAIKILKDVIIAREEVAHTLTENRVLQRC 32493

Query: 206 RHPFLT 211
+HPFLT
Sbjct: 32494 KHPFLT 32511

Score = 190 (87.6 bits), Expect = 5.2e-165, Sum P(7) = 5.2e-165
Identities = 36/45 (80%), Positives = 37/45 (82%), Frame = +2

Query: 276 KLENLMLDKDGHKITDFGLCKEGIKDGATMKTFCGTPEYLAPEV 320
KLENL+LDKDGHIKI DFGLCKE I G TFCGTPEYLAPEV
Sbjct: 33509 KLENLLLDKDGHIKIADFGGLCKEISFGDKTSTFCGTPEYLAPEV 33643

Score = 188 (86.7 bits), Expect = 5.2e-165, Sum P(7) = 5.2e-165
Identities = 37/57 (64%), Positives = 42/57 (73%), Frame = +3

Query: 209 FLTALKYSFQTHDRLCFVMEYANGGELFFHLSRERVFSERARFYGAIEVSALDYLH 265
+ LKYSFQ LCFVM++ANGGELF H+ + FSE RARFYGAIEV AL YLH
Sbjct: 32667 YFQELKYSFQEQHYLCFVMQFANGGELFTHVRKCGTFSEPRARFYGAIEVLALGYLH 32837

Score = 166 (76.5 bits), Expect = 5.2e-165, Sum P(7) = 5.2e-165
Identities = 29/59 (49%), Positives = 42/59 (71%), Frame = +1

Query: 53 NNFSVAQCQLMKTERPRPNTFIIRCLQWTTVIERTFHVETPEEREWEATAIQTVDGLK 111
+ F++ Q M E+PRPN F++RCLQWTTVIERTF+ E+ E R+ W AI++++ K
Sbjct: 31846 STFAIFYFQTMLFEKPRPNMFMVRCLQWTTVIERTFYAESAEVRQRWIHAIESISKKYK 32022

Score = 134 (61.8 bits), Expect = 5.2e-167, Sum P(8) = 5.2e-167
Identities = 24/33 (72%), Positives = 30/33 (90%), Frame = +3

Query: 210 LTALKYSFQTHDRLCFVMEYANGGELFFHLSRE 242
L LKYSFQT+DRLCFVME+A GG+L++HL+RE
Sbjct: 33156 LQELKYSFQTNDRLCFVMEFAIGGDLYYHLNRE 33254

Fig. 25



Fig. 26A

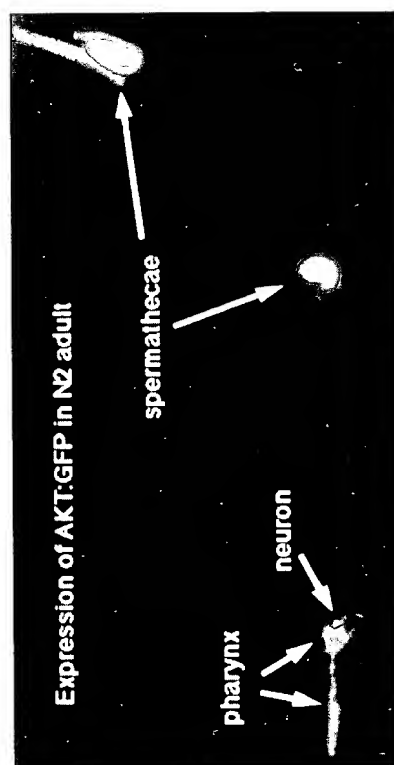


Fig. 26B

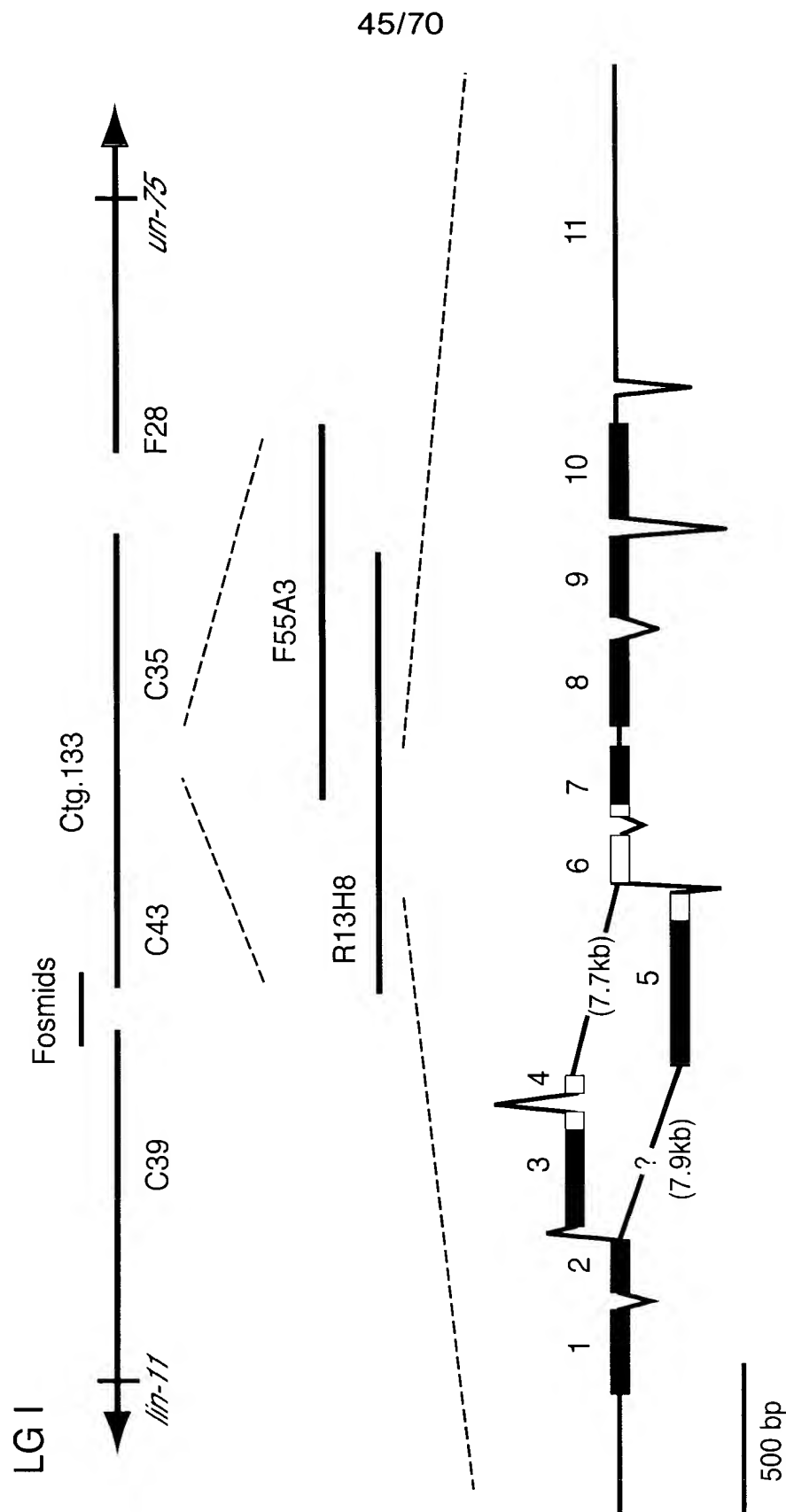


Fig. 27



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	1	15	16	30	31	45	46	60	
1 ZK84.6	-MNSVFTIIFVLCAL	QVAASFRQSFG---	P	SMSEESASMQLLREL	QH--	NMMESAHRPMP			54
2 ZK75.1	-MFSFFT-YFLLSAL	LLSASCRQ-----	P	SMDT-SKADRILREI	E----	METELLENQLS			47
3 ZK1251.2	----MPPIILVFFLV	LIPASQY-----	P	FSLE-SLNDQIINEE	VI--	EYMLENSIRSS			47
4 C06E2	--MIVTLIVFLVIGL	QMAHLSQVSGNNENG		FLNP-FDLSQWSEEI	LHRQYHHHHHHHHGN				57
5 ZK75.2	----MNAIIFCLLFT	TVTATYEVF-----	G	KGIEHRNEHLIINQL	D---	IIPVESTPTPN			48
6 ZK75.3	MKLSVVLALFIIFQL	GAASLMRN-----	W	MDFEKELEHDYDDS	E---	IGFHNIHSLMA			51
7 C17C3	-----	-----		-----MKLLHI	F---	IIFLLFQSCSN			18
8 F13B12	-----	-----		---MYWFRQVYRPS	FF--	FGFLAILLLSS			50
9 INSULIN	-----	-----		-----MA	LWMRLPLLLALLALW				17
CONSENSUS	-----	-----		-----	-----	-----			

	61	75	76	90	91	105	106	120	
1 ZK84.6	RARRVPAPGETRACG	RKLISLVMVAVCGD-L	CN-----						85
2 ZK75.1	RARRVPA-GEVRACG	RRLLLFVWSTCGE-P	CT-----						77
3 ZK1251.2	RTRRVPDEKKIYRCG	RRIHSYVFAVCGK-A	CE-----						78
4 C06E2	RARRTLETEKIYRCG	RKLYTDVLSACNG-P	CE-----						88
5 ZK75.2	RASRVQK----RLCG	RRLILFMLATCG--E	CD-----						74
6 ZK75.3	RSRRGDK---VKICG	TKVLKMMVMVCGG-E	CS-----						79
7 C17C3	KMCQYSK-KKYKICG	VRALKHMKVYCTR-G	MT-----						48
8 F13B12	PTPSDAS---IRLCG	SRLTTTLLAVCRNQL	CTGLTAFKRSADQSY	APTTRDLFHIHHQQ-					80
9 INSULIN	GPDPAAAFVNQHLCG	SHLVEALYLVCGERG	FFYTPKTRREAEDLQ	VGQVELGGGPGAGSL					77
CONSENSUS	-----CG	-----C-----	-----	-----	-----	-----	-----	-----	

B CHAIN

C PEPTIDE

	121	135	136	150	151	165	166	180
1 ZK84.6	-----PQEGKDIA	TECCGNQCSDDYIRS	ACCP-----			112		
2 ZK75.1	-----PQEDMDIA	TVCCTTQCTPSYIKQ	ACCPEK---			106		
3 ZK1251.2	-----SNTEVNIA	SKCCRECTDDFIRK	QCCP-----			105		
4 C06E2	-----PGTEQDLS	KLCCGNQCTFVEIRK	ACCADKL--			118		
5 ZK75.2	-----TDSSDLS	HICCIKQCDVQDIIR	VCCPNSFRK			106		
6 ZK75.3	-----S-TNENIA	TECCEKMCTMEDITT	KCCPSR---			107		
7 C17C3	-----R-DYGKLL	VTCCSKGCNAIDIQR	ICL-----			73		
8 F13B12	-----KRGGIA	TECCEKRCSFAYLKT	FCCNQDDN-			109		
9 INSULIN	QPLALEGSLQKRGIV	EQCCTSICSLYQLEN	YCN-----			110		
CONSENSUS	-----	-----CC---C-----	-----C-----	-----	-----	-----	-----	-----

A CHAIN

Fig. 28



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Zk75-1	ACGRRLLLV	WSTCGEPCTx	xxQEDMDIAT	VCCTTQCTPS	YTKQAC46
Zk84-6	Acgrklisl	maVcgdlcnx	xxqegkqiat	eccgnqcdd	Ylrsac46
Zk1251-2	RCGRRHSYV	FAVCGKACEx	xxSTEVENIAS	KCCREECTDD	FIRKQC46
C06e2	RCGRKLYTDV	LSACNGPCEX	xxGTEQDISK	LCCGNQCTFV	EIRKAC46
Zk75-3	TCGTRV LKMV	MVMCGGECsx	xxSTNENIAT	ECCEKMCTME	DIITTKC46
Zk75-2	lcgrrlilfm	latcgecdtx	xxDSSEDLISH	ICCIKqcdvq	dilrv46
Ins-Human	LCGSHLVEAL	YLVCGERGfx	xxLQKRGIVE	QCCTSIcsLY	QLENYC46
Ins-Rabbit	lcgshlveal	ylvcgergfx	xxtpksgive	ccctsiCSly	qlenyc46
Ins1-Xenopus	lcgshlveal	ylvcgdrgfx	xxkmkrgive	ccchstcslf	qlenyc46
Ins2-Xenopus	lcgshlveal	ylvcgdrgfx	xxkmkrgive	ccchstcslf	qlenyc46
Ins-Alligator	lcgshlvdal	ylvcgergfx	xxspkggive	ccchntcsly	qlenyc46
Ins-Elephantfish	lcgshlvdal	ylvcgergfx	xxpkqigive	ccchntcsly	qlenyc46
Igf1-Bovine	LCGAELVDAL	QFVCGDRGFx	xxAPQTGIVD	ECCFRSCDLR	RLEMYC46
Igf1-Dog	lcgaellvdal	qfvcgdrgfx	xxapqgtgvd	eccfrscdlr	rlemy46
Igf2-Horse	lcggellvdtl	qfvcgdrgfx	xxrrsrgive	eccfrscdlr	lletyc46
Igf2-Human	LCGGE LVDTL	QFVCGDRGFx	xxRRSRGIVE	ECCFRSCDLA	LLETYC46
Ilp-Amphioxus	LCGSTLADVL	SFVCGNRGYx	xxRRRRRGIVE	ECCYNVCDYS	QLESYC46
Lirp-Locust	YCGEKL SNAL	KLVCRCGNYNx	xxRRTRGVFD	ECCRKSCSIS	ELQTYC46
Bxa4-Bommo	YCGRHLARTL	ADICWEAGVx	xxRGKRGIVD	ECCLRPCSDV	VLLSYC46
Bxb1-Bommo	YCGRHLADTL	ADICFGVEKx	xxRGKRGVVD	ECCFRPCTLD	VLLSYC46
Bxrpa-Hornworm	icgrhlartl	adlcpnveyx	xxgkragvad	ccvnsctmd	vllsyc46
Bxa1-Silkworm	ycgrrlatml	sfvcnqyqx	xxgkrqgiae	eccnkpcten	ellgyc46
Bxa2-Silkworm	YCGRRLATML	LYVCDNQYQx	xxGKRQGIVE	ECCNKPC TEN	ELLGYC46
Bax3-Silkworm	ycgrrlaiml	syicnqylx	xxgkrqgiae	eccnkpcted	ellgyc46
F13b12	LCGSRLTTTL	LAVCRNQLCx	xxQKRGGIAT	ECCEKRC SFA	YLKTFC46
Mpi3-Seasnail	LCGSTLANMV	QWFCSTYTTx	xxESRPSIVC	ECCFNQCTVQ	ELLAYC46
Relaxin-Human	LCGRELVRAQ	IAICGMSTWx	xxRPYVALFE	KCC LIGCTKR	SLAKYC46
Rlf-Human	lcghhlvral	vrvcggprwx	xxaaatnpar	Ycc lsgctqq	dllt1c46

Fig. 29

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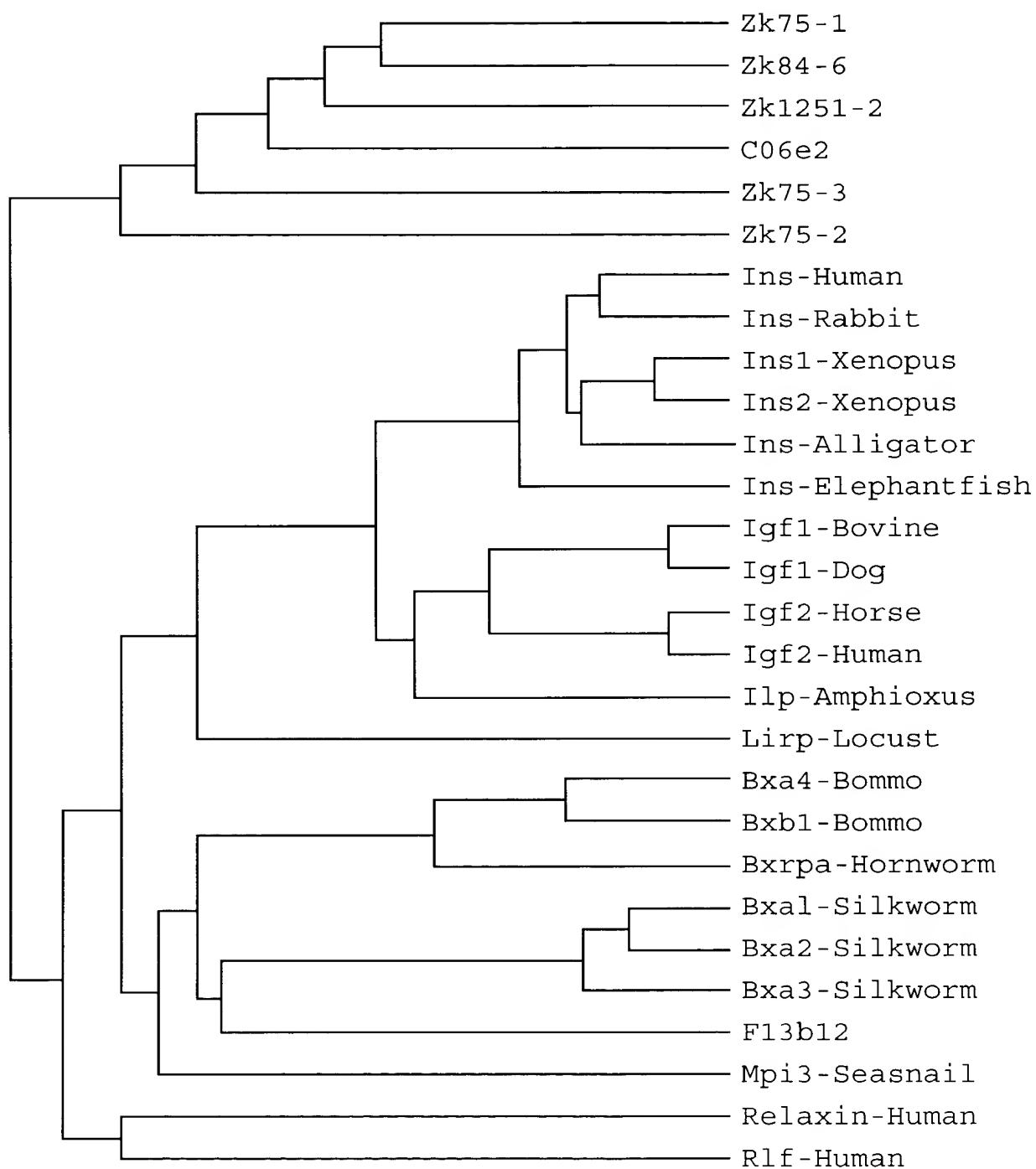


Fig. 30

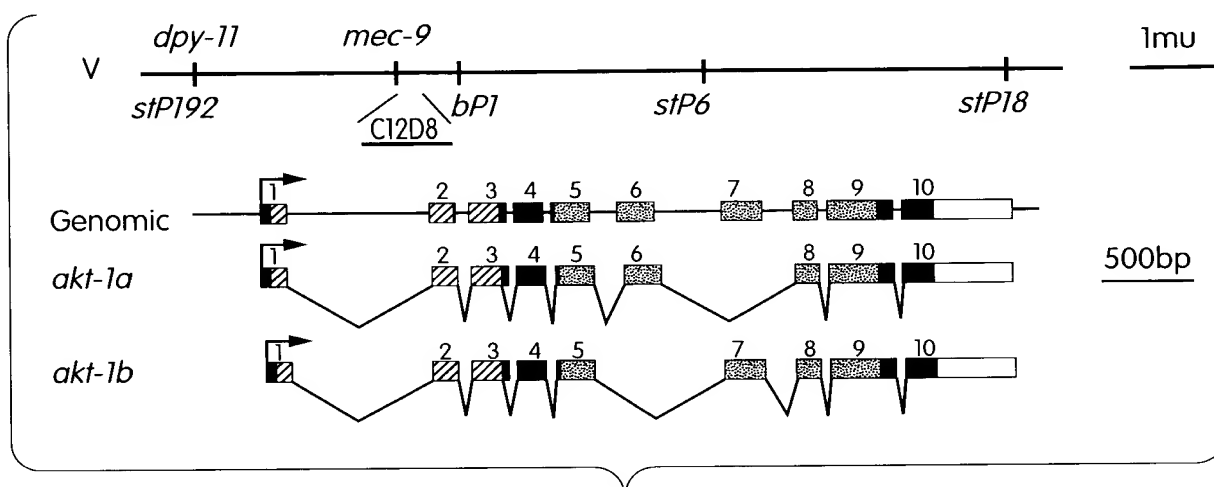


Fig. 31

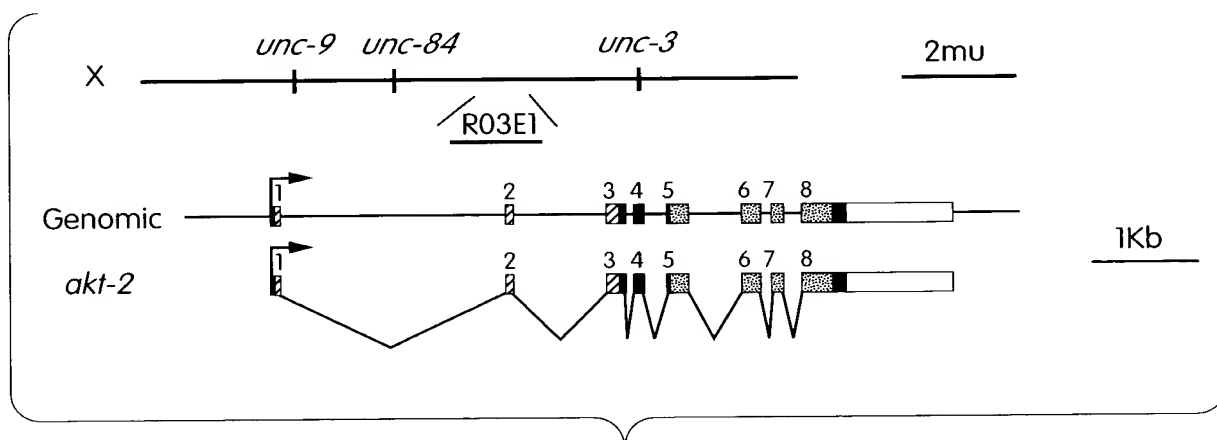


Fig. 32

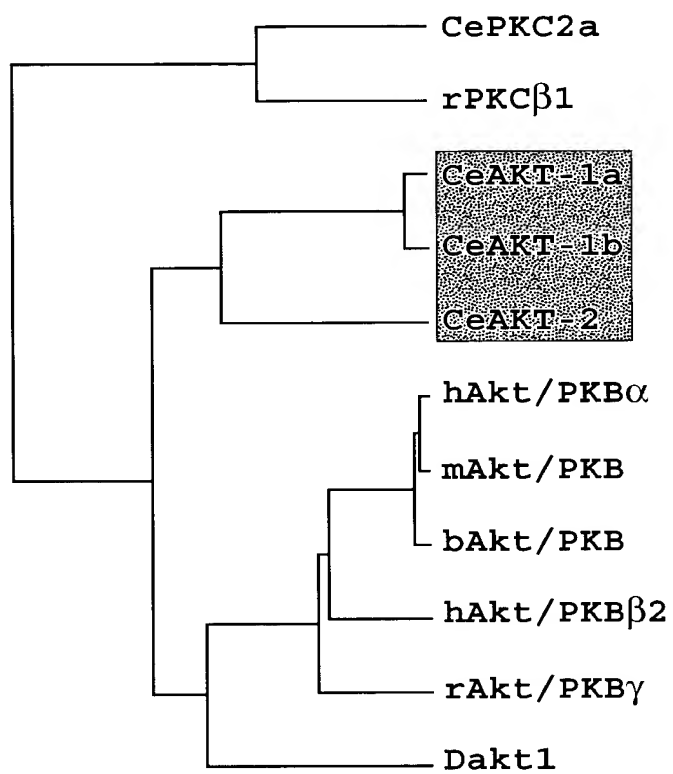


Fig. 33



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AKT-1a MSMTSLSTKSRR--QEDVVIEGWLHKKGEHIRNWRPPYFMFNDGALLGFRAPKKEGQPFPEPL
AKT-1b
AKT-2 M..ENAHLOK..I..S.. IL R T S D L
hAkt/PKBa MSDVAI K R Y KT LLK TFI YKER QDVDQREA

AKT-1a NDFMIKDAATMLFEKPRPNMFMVRCLQWTTVIERTFYAESAEVRORWIHALESIS--KKYKGTN
AKT-1b
AKT-2 N R VCLD I D DF E QAV SHNR ENA
hAkt/PKBa N SVAQCQL KT R T II HV TP E EE TT QTVADGL KOE--
mg144 T

AKT-1a ANPQEELMETNQPKIDEDSEFAGAAHAIMGQPSSGHGDNC SIDFRASMISIADTSEAAKRDKI
AKT-1b
AKT-2 G.TSMQEED..GN.SGES.VNM-----DAT.TRS.....ESTVMN.DEPE.VPRKNTV
hAkt/PKBa -----E.EMD-----R.GSPS..SGAE-----EMEV.L.KPKHRV

AKT-1a TMEDFDLKVLEKGTGKVLCKEKRTOKLYATKFLKDDVIAREEVAHTLTENRVLQCKHPF
AKT-1b
AKT-2 ..D. Q R SSD IR EMVVD S YA V
hAkt/PKBa ..NE EY L V A GRY M E V KD NSR

AKT-1a LTELKYSFOEQHYLCFVMOFANGGELFTHVRK---CGTFSEPRARFYGAIEIVLALCYLH RC
AKT-1b TND R E T D VV LNREVQMNKEG S AN
AKT-2 L A YHI E LQR ---K A T S I HR
hAkt/PKBa A THDR EY F LSRE---RV D S D SEK

AKT-1a DIVYRDMKLENLLLDKDGHIKIADFGLCKEEISFGDKTSTFCGTPEYLAPEVLDDHDYGRVDW
AKT-1b S L
AKT-2 N R T KY IE I D S
hAkt/PKBa NV L M T G KD ATK E N A

AKT-1a WGVGVVMYEMMCGRLPFYSKDHNKLFELIMAGDLREPSKLSQEARTLLTGLLVKDPTQRLGGGP
AKT-1b
AKT-2 SA ENG TTC K NR P V S ERV AK A
hAkt/PKBa L NQ E LMEEL RT GP KS S K K S

AKT-1a EDALEICRADFFERTVDWEATYRKEIEPPYKPNVQSETDTSYFDN-EFTSQPVQLTPPSRSGALA
AKT-1b
AKT-2 D R VS E KD.....L..V...F...M.....F..RVRYV.ILLKV-----E.I
hAkt/PKBa K MOHR AGIV.QHV.E.KLS..F..Q.T....R...E-...A.MITI...DQDDSME

AKT-1a TVDEQEEMQSNFTQFSFHNVMGSINRIHEASEDNEDYDMGZ
AKT-1b
AKT-2
hAkt/PKBa C---S.RRPH.P...YSASSTA

Fig. 34



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cataaaaatccagtaaatggtaaaattttcaatttcagatccatctcgatggaggatctcacaccaactaacacgtcgctcgacaccacaactac
taacaatgacacgacatcggtatcgatgaagcgccgccaacggtaggaactagtttctagacgaacatcggaatgcggcttaaagttcgggtgcac
ttatcaaactagaccggttttttagacctctttcaaagcggggaactgcaatacactttttgaacctaaacctagatttttgggtgttctaaat
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aatccaggagtggatttctcgttcccagaaggatttccagaggaagcgtcggaattatcgcaag

Fig. 35A



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at tt t t g g t a g g t t g a c a t g a a a c t t t a a a a c t g a a t a c g t a a t t t t c a a c t t a c a g g t g c g c g a c c c g a g t a c c c g t a t c a c c a g t c a a g a a c t
t a t g g c t c a c a a g t t t t t g a a a c g t t g a c t g g g t g a a c a t t g c a a a t a t c a a g c c a c c a g t c c t g c a c g c c t a c a t t c c a g c c a c a t t t g g c g
a g c c g g a g t a c t a c t a c a c a t t g g g c c t g t c g a g c c g g a c t t g a t g a t c g t g c c t t g t t c c g t t t g a t g a a t t t g g g a a a t g a t g c t a g c g c a
t c a c a g c c a t c a a c g t g a g t t t g a a g c a t t t t t t c t t g c a t t a a a a g t t t t a c c t t g c a c t g a c c a a a a t t t a t t g a a a c t a t t a a t t a t t t g a
t t c t g a t t a a c a a t g a c c a a a g a t t t g a a c t g a c a a a g t g c a a a t t t g c a c c g a c c a a a a a c a g t t t g c a c t g a c c a c c t c t t c a t t t g c a c t
g a c c a c c t c t t c a t t t g c a c t g a c c a a c t t t t c a t t t g c a c t g a c c a t c t c t t c a t t t g c a c t g a c c a a c t t t t c a t t t g c a a t t c t g g c a a t g a
t t c t t t t g c a t c t a c t g a t c a a a a a t t g a t t c a a a t c a a t t a a t t t t c t t t g a c a g t a c t a t g c c t t a t t c a a g g a g a t g c t g a t c t g a a a a t t c
t c a a t a g t t g a t a a a a a t t a c t a a c c c c t t a g a a a g t t t c a g a c c g t c t a a c g t g g a a c a t c g c g g a g a c c c a t t t g t t c g g a a a t t g c a c c g t
g a g t g a t t t g c a c c t a a t t g g t t a t t t t a a t a a t c a t t a a a t t a t a g a c g c c a a t t c g g a a g c c g a a a a g a a c c g c c c g c a c g t g c g c a g a
a g c t c g a a g a g a c a c g t g t c a a a a c c c a t t c c a c a t c t t c a c c a a c a a c t c g c t c a t t t t g a a a c a a g g a t a t t t g g a a a a g a a g c g a g g a t t g
t t t g c c a g a c g c c a a t g t t c c t g t t g a c c g a a g a c c g c a t c t c t t g t a c a t t g a t g t g c c g a a t c t t g t g c t c a a a g g a g a g t a c c a t g g a c
g c c g t g c a t g c a g g t g g a g c t a a a a a c t c g g g a a c t t t c t t a t a c a t a c g g t a g g t c a g a a t a a t c a t a g c t g t c t a t c t a t t a t a g t a c t c
a a t g a a t c t g a a a a t t c a a a t t t t c a g c c c a a c c g c t c t a c t a c t t g t t t g a t c t c g a a a a g a a g c a g a t g a g t g g t g t a a g g c t a t c a a t g
a t g t t c g c a a g c g g t a c t c g g t g a c t a t c g a a a a g a c t t t t a a c t c t g c g a t c g g t g a c g g a a c a t t t g g c a g c a t t t a t g g a a a g a a a a g t c c
a g a a a g g t a t g a a t t a c t g g a a g c c c c c t c a c t g a g t t t c c a g c a a g t t c a g a g t t t t t a t t g g a a t t t t g c c a a t t t t c a t t a g a c t t t a
g a g c c t a t t g c t a t t t t g t g g a c a g g t t t a a c a t t t t c a a a a a a a a t t g a g a a t g t c t g a a a a a t t t g g a g t g t g a c a g t t t t c t g a a t t t
t g a a a a t t c t g t t c t c a a a a t t g g a t t t t a c a g a c t t g t t t c g a g a t t t c a t a a t c c t t c a a a g a a t a t a g a a t a t t t g t g t t c a a c t t t t c
t t g t c a a a a t a t t t t t t t g g a c a a t c t a g a t t c t g g a a a t t t t c a a a a a a g a t a a t c t c t a a a c a a a c t a a a t t c a a a t g t t c t a a a g g t
t c t t t a t t t t c c a t g c a a c t c t a a a a t c t t c c c g t a t a t t t t t t g g a a a g t c t t a t g a t g t t a g a c g g t t t a a a t t t t t g a t g a t t t a a a t t
t t t t a g g g t g g t c t a t a a t t t t g g a c c a c c c t g t a t a a t t a t g g a c c a c c a t g t a c a c t t a t a g a c c a c c c a g t a a c a a g c a t t t t t g g a c c a c
c a c g c a a a t c t t a t t a t t a t g g a c c a c c c a a c t t a g a a c a c c t t c a a t a c t t c t t t t c t g t t c a a a a a t g a t c a a c t t g c t g a a a a a a a t t t
t t t g t a g g a a a t g a t c g c t g a a c a g a g c g c t g c c c g c a a a c a a g a a a a g g a g a g a a a a g c g c t a a a a g c c g a g c a a g t g a g c a a g a a g c
t t t c a a t g c a a a t g g a c a a g a a g t c g c c t t g a a g g c t c a c c t c c c t t c t a c t c c c c a c a a a t c a c c a t c a a c a a a t c a c a c t t t t g t a t c a t t
t t g c g t c c

Fig. 35B



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MEDLTPTNTSLDTTTTNNDTTS DREAAPTTLNLTPTASESENSLSPVTAEDLIAKSIKEGCPKRTSND FMFLQSMGEG
AYSQVFRCREVATDAMFAVKVLQKSYLNRHQMDAIIREKNILTYLSQECGGHPFVTQLYTHFHDQARIYFVIGLV
ENGD LGESLCHFGSFDMLTSKFFASEILTGLQFLHDNKIVHRDMKPDNVL IQKDGHILITDFGSAQAFGGLQLSQEGFT
DANQASSRSSDSGSPPPTRFYSD EEEENTARRTT FVG TALYVSP EMLADGDVGPQTDI WGLGCILFQCLAGQPPFRAV
NQYHLLKRIQELDFS FPEGFP EEEASEIIAKILVRDPSTRITSQELMAHKFFENVDWVN IANIKPPVLHAYIPATFGEP
EYYSNIGPVEPGLDDRALFRLMNLGNDASASQPSTPSNVEHRGDPFVSEIAPRANSEAEKNRAARAQKLEEQRVK
NPFHIFTNNSLILKQGYLEKKRGLFARRRMFLLTEGPHLLYIDV PNLVLKGEVPWTPCMQVELKNSGTFFIHTPNR
VYYLFDLEKKADEWCKAINDVRKRYSVTIEKTFNSAMRDGTFGSIYGKKKSRKEMMREQKALRRKQEKEKKAL
KAEQVSKKLSMQMDKKSP

Fig. 36

MEDLTPTNTSLDTTTTNNDTTS DREAAPTTLNLTPTASESENSLSPVTAEDLIAKSIKEGCPKRTSND FMFLQSMGEG
AYSQVFRCREVATDAMFAVKVLQKSYLNRHQMDAIIREKNILTYLSQECGGHPFVTQLYTHFHDQARIYFVIGLV
ENGD LGESLCHFGSFDMLTSKFFASEILTGLQFLHDNKIVHRDMKPDNVL IQKDGHILITDFGSAQAFGGLQLSQEGFT
DANQASSRSSDSGSPPPTRFYSD EEEVPEENTARRTT FVG TALYVSP EMLADGDVGPQTDI WGLGCILFQCLAGQPPFR
AVNQYHLLKRIQELDFS FPEGFP EEEASEIIAKILVRDPSTRITSQELMAHKFFENVDWVN IANIKPPVLHAYIPATF
GEPEYYSNIGPVEPGLDDRALFRLMNLGNDASASQPSTFRPSNVEHRGDPFVSEIAPRANSEAEKNRAARAQKLEE
QRVKNPFHIFTNNSLILKQGYLEKKRGLFARRRMFLLTEGPHLLYIDV PNLVLKGEVPWTPCMQVELKNSGTFFIH
TPNRVYYLFDLEKKADEWCKAINDVRKRYSVTIEKTFNSAMRDGTFGSIYGKKKSRKEMMREQKALRRKQEKEE
KKALKAEQVSKKLSMQMDKKSP

Fig. 37



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Fig. 38A



Fig. 38B



Fig. 38C



Fig. 38D



Fig. 38E



Fig. 38F



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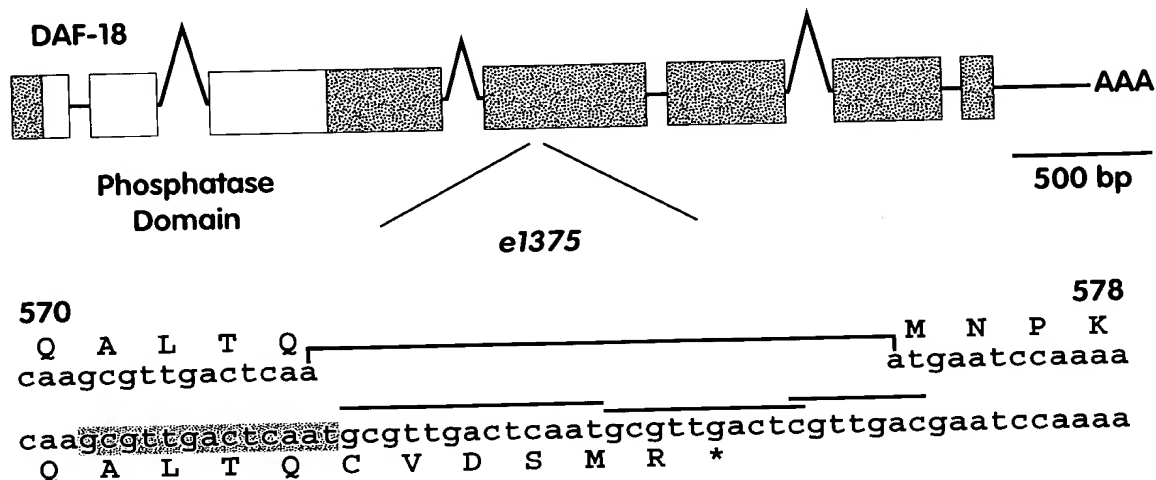


Fig. 39A

DAF-18	48	IFRTAVSSNR	CRTEYQNI	DLDCAYITDRI	AI	GYP	PAIGIE	ANFRNSKVQT
PTEN	4	IIKEIVSRNK	RRYQEDGFDL	DLTYIYPNFI	AMGFPAERLE			GVYRNNIDDV
DAF-18	98	QOELLTRRHGK	GNVKVFNLRG	GYYYDADNED	GNVICFDMTD	HHPPSLEELMA		
PTEN	54	VRFLDSKH.K	NHYKIYNLCA	ERHYDTAKEN	CRVAQYPFED	HNPPQLELIK		
DAF-18	148	PEGREAKEWL	EADDKHVI	AVHCKAGKGRTG	VMICALIYI	NFYPSPROIL		
PTEN	103	PECEDLDQWL	SEDDNHVAAT	HCKAGKGRTG	VMICAYELHR	GKFLKAQEAL		
DAF-18	198	DYYSIIRTKN	NKGVITPSQR	RYIYYHKL	ERELNMLPLR	MLIGVYVER		
PTEN	153	DFYGEVRI	RDKKGVTIPSQR	RYVYYSYLL	KNHLDYRPVA	LLFHKMMFET		
DAF-18	248	PPKTWGGGSK	IKVEVNGNST	ILFKPD..PL	IISKSNHQRE	RATWLNNDT		
PTEN	203	IPMFSGGTCN	PQFVVCQLKV	KIYSSNSGPT	RREDKFMFE	FPQPLPVC		

Fig. 39B



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DAF-18 Protein

MVTPPPDVSTSRSMARDLQENPNRQGEPRVSEPYHNSIVERIRHIFRTAVSSNRCTEYQNIDLDCAIYITDRIIAIG
YPATGIEANFRNSKVQTQQFLTRRHGKGNKVFNLRGYYDADNFDGNVICFDMTDHHPSPSLELMAPFCREAKEWLEAD
DKHVIADVHCKAGKRTGVMICALLIYINFPSPRQIILDYYSIRTKNKGVTIPSORRYIYYHKLRERELNYLPLRMQL
IGYVERPPKWTGGGSKIKVEVGNGSTILFKPDPDLIISSNHQERATWLNCCDTPNEFDTGEQKYHGFVSKRAYCFMVP
EDAPVFVEGDVRIDIREIGFLKKFSDGKIGHVWENTMFACDGLNGGHFEYVDKTPYIGDDTSIGRKNMRRNETPMRK
IDPETGNEFESPWQIVNPPGLEKHITEEQAMENYTNYGMIPPRYTISKILHEKHEKGIKDDYNDRKLPMDKSYTESGK
SGDIRGVGGPFEIPYKAEHVLTPVYEMDRALKSKDLNNGMKLHVLRVCVTRDSKMMKSEVFGNLAHNESRRRLQA
LTQMNPKWRPEPCAFGSKGAEMHYPPSVRYSSNDGKYNACSENLSVDFFEHRNIAVLNRYCRYFYKQORSTSRSRYPKRF
RYCPLIKKHFIYPADTDDVDENGQPFHSPHEHYIKEQEKIDAEKAAKGIENGTGPSTSGSSAPGTIKKTEASQSDKVKPAT
EDELPPARLPDNRFRFPVGVDFENPEEESCEHKTVESIAGFEPLHLEHSHYHNTAGNMLRQDYHTDSEVKIAEQEAK
AFVDQLLNGQGVLEFQKQFVPSDNSFADYVTGQAEVFKAQIALLEQSEDFQVQANAEVVDLEHTLGEAFERFGHVVE
ESNGSSKNPKALKTREQMVKETGKDTQKTRNHVLLHLEANHRVQIERRETCPHELHPEDKIPRIAHFSENSFSDSNFDQAI
YL

Fig. 40A

1 ttcagggtac atctactaac ccccaatgggt tactctctct cagatgtgc caagcacatc
61 gaccaggtcg atggctcgtg accttaaga gaatccaaac cgaacacctg gtgaaccacg
121 tgggtctgaa cgtatccaca atccaatcgt cgagcggatt cgcatattt ttccgacggc
181 tgtatcttcc aatcgttctc gcaccgagta ccaaatatc cactagatt gtgcataat
241 cagagaccga atcatagcta tgggttatcc agcaacagga agcaaggca atttccgtaa
301 ctcaaaagt ctcaactcaac aatctctgac caggcggac tgcggataac atgttaattg
361 gtttaacctg cgcggtggat actactacg actcgaata tggctccgt ttgacagaga
421 ctcgatatg actgatcat cccgctgaag cagacgataa acatgaata cgtgtacact gtaagctgg
481 ggttaaggaa tggcttgaag tggatgtgc tctctctatc tctctctatc tacatcaact tctatccgag
541 aaaggccgt accggtgta tgaatgtgc actactcaat aatctctaca aaaaacaaca aggtgtcac
601 cccacgacaa atctctgact actactcaat acatttacta ctaccataag ctctcgtgac gtgagctcaa
661 aattccatca caacgacgct agttgattgg agtttctcaa aagtttctg gacgggaaga ttggtcattg
721 ctatttaccg ttgagaaatgc aagtggaggt tggaaatggc tggactcaac ggtggacatt cggagtacgt
781 ggtggtgggt tcaagataaa aatcaaatca aatcgaatga tccagcagag agcgaacaa aggtgaacaa
841 ggtatctctc ataatctcca tgcacacgg tgcacagaaa agagcaaaa tgcgtcagta ttgctcgaag gagatgttgc
901 ctgtgatcgc cctaacgaat tgcagacgg tgcagagaa agactcaac aggttctgc gacgggaaga ttggtcattg
961 gagagcatc cgcgaaatcg gatctctcaa atgttctgc aggttctgc gacgggaaga ttggtcattg
1021 tatagacat tgggttcaat acaatgttctc cgcgaaatcg gatctctcaa atgttctgc aggttctgc gacgggaaga
1081 tgggttcaat acaatgttctc cgcgaaatcg gatctctcaa atgttctgc aggttctgc gacgggaaga
1141 agacaaaact cgcgaaatcg gatctctcaa atgttctgc aggttctgc gacgggaaga
1201 aagaaaatgaa acgctgatgc gaaatctga tccagaaat tccagaaat ggaatgaat ggaatgaat
1261 gggcaataa gtgaatcttc ttgacttgc tgaatctga tccagaaat tccagaaat ggaatgaat ggaatgaat
1321 aatlatacc aatlatgga tcaagtgatg tcaagtgatg atataatgc cgaatgaat atcagcaga cgaatgaat
1381 aagcatgaa aaggtatcg aaggtatcg aaggtatcg atataatgc cgaatgaat atcagcaga cgaatgaat
1441 caaatcatc aaggtatcg aaggtatcg aaggtatcg atataatgc cgaatgaat atcagcaga cgaatgaat
1501 gataccatc aaggtatcg aaggtatcg aaggtatcg atataatgc cgaatgaat atcagcaga cgaatgaat
1561 attgaagagt aaagatctta aaggtatcg aaggtatcg atataatgc cgaatgaat atcagcaga cgaatgaat
1621 tactcgtgat tcaaaaatga cggaggtctc aaggtatcg aaggtatcg atataatgc cgaatgaat atcagcaga
1681 tgaatcgaca cggaggtctc aaggtatcg aaggtatcg atataatgc cgaatgaat atcagcaga cgaatgaat
1741 ggtggttcc ggtatccaaag ggtatccaaag ggtatccaaag aaggtatcg aaggtatcg atataatgc cgaatgaat
1801 caatgatgga agtatataatg agtatataatg agtatataatg aaggtatcg aaggtatcg atataatgc cgaatgaat
1861 cagaaatatt gccgttctta atccagata aatccagata aatccagata aatccagata aatccagata aatccagata
1921 tgaagccgt accgatgat accgatgat accgatgat aatccagata aatccagata aatccagata aatccagata
1981 tccagctgat accgatgat accgatgat accgatgat aatccagata aatccagata aatccagata aatccagata
2041 ttacatlaaa acttcaggat acttcaggat acttcaggat aatccagata aatccagata aatccagata aatccagata
2101 tggaccagg acttcaggat acttcaggat acttcaggat aatccagata aatccagata aatccagata aatccagata
2161 atccgacaag ttccagtcg ttccagtcg ttccagtcg aatccagata aatccagata aatccagata aatccagata
2221 tgtcggaaga ttccagtcg ttccagtcg ttccagtcg aatccagata aatccagata aatccagata aatccagata
2281 acacaaaacc gtagagtcga gtagagtcga gtagagtcga aatccagata aatccagata aatccagata aatccagata
2341 ataccatcca aatccagtcg aatccagtcg aatccagtcg aatccagata aatccagata aatccagata aatccagata
2401 gaaaatagct gaacaaaggg ttatgaagc ttatgaagc ttatgaagc ttatgaagc
2461 attacaagag ttatgaagc ttatgaagc ttatgaagc ttatgaagc ttatgaagc ttatgaagc ttatgaagc

Fig. 40B-1



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2521 aaccggacag gccgaagttt ttaaagcaca gattgcgtaa ctggagcagt cggaggattt
2581 tcaacgagtt caagcgaatg cagaggaagt cgatcttgaa cacactcttg gtgaagcgtt
2641 tgagcgattc gggcacgttg tagaagaatc gaatggttct tctaaaaatc caaaagccct
2701 gaaaactcga gaacaaatgg tgaagaaaac tggcaaaagc actcagaaga ccgcaatca
2761 tgtgcttcta catttgaag ctaatcatcg tgtgcaaatc tggcaaaagc gagctcgtg aacgtgccc
2821 ggagctacat ttgatcaag aaatcccaag aattgctcat tttccgaaa acagcttctc
2881 ggattcgaat ttgatcaag ctatttattt gtaaacctaa acaaaactt ttagaagatt
2941 ttcttcttac tgacctcca attttcagat aatttcaatg acttaagttt tctttcaaa
3001 gtatcattca ctttctgtat agtgttttgt ttttaacaa actatgttc gattattttg
3061 tatattcata ttatagctct caacttccc atttccacg tataatgta tattttgccg
3121 ggtgaaaaat agcaattccc tatgaatgta tcccccca tctgtttct tactcagaaa
3181 ttgtaattca cattgcgggt catcactaat cctatgggt ttaacacaat tctcccataa
3241 atcaattgta cttaccaatt ttttgtttaa ttatttagat ttgtaacatt gaaattggtg
3301 ataa

Fig. 40B-2

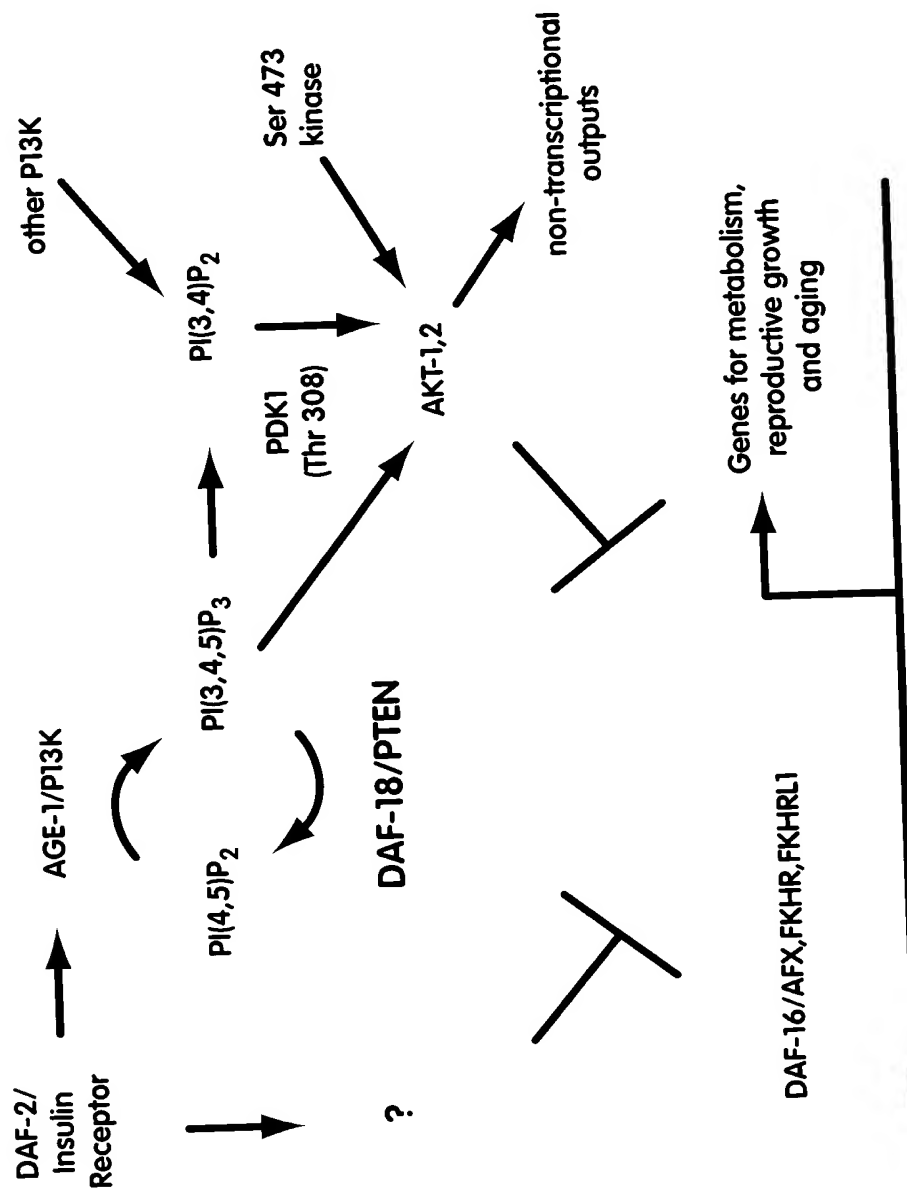


Fig. 41



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ttta
attaccaagtttgaggtagcatgtgctctcttcaatcat atg gat tcg ttg ttt cag atg gca tcc gca
M K F Q Y Y S K K A A gct gct gga aag aca atg tct aat agt gtc tcc
atg tcc agt gac aat cgc atg gag gat ttt aaa cgt cgt ttt cgt cga agt gga tgc tta
M S S D N R M E D F K K R R F R R R S G S L
gga att cca ttt gtc cca gaa gaa gat gtt aaa cca ctc ttc aca cca act cgt act gtt
G I P F V P E E D V K Q L F T P T R T V
cgt cga gaa gca tst att cgc gaa gaa ggg gat gag gaa gaa gta caa att ctc aca ata
R R E A S I R E G G D E E E G V R I L T I
att gtc aag tca agt cgt gtt gtt tgc gag gat atc tca aaa atg att gca aac ctc cct gat
I V K S S R V V S E D I S K M I A N L P D
cac act cgt atc aaa cat ttg gag act cgt gag act cca gat gga agt tcc aaa act atg
H T R I K H L E T R D S Q C D G S S K T M
gat gtt ctt cta gag att gag ctc ttt cat tat gga aaa caa gaa gca atg gat ctt atg
D V L L E I E L F H Y G K Q E A M D L M
aga ctt aat ggg ctt gat gtt cat gag gtg tca tgc act att cgt cca act gca ata aaa
R L N G L D V H S S T I R P T A I K
gag caa tat aca gag cct gga tct gat gat gcg aca acc ggt tct gaa tgg ttt cca aaa
E Q Y T E P G S D D A T T T G S E W F P K
agt att tat gat ttg gat att tgt gca aaa aga gtg att atg tat gga gca ggg ctg gac
S I Y D L D I C A K R V I M Y G A G L D
gct gat cat cct ggt ttc aaa gat acc gag tat cgt caa cgt cga atg atg ttt gct gaa
A D H P G F K D T E Y R Q R R M F A E
ctg gcg ctc aat tac aaa cac ggt gag cca att ccg cga acc gaa tat aca tca tcc gaa
L A L N Y K H G E P I P R T E Y T S S E
cgg aaa act tgg gga att ata tat aga aaa ttg aga gaa ttg cac aaa aag cac gca tgc
R K T W G I I Y R K L R E L H K K A C

Fig. 42-1

aag cag ttt ctt gat aac ttt gag cta ctg gag aga cat tgt gga tac tcg gaa aat aat
K Q F L D N F E L L L E R H C G Y S E N N

att ccg caa cta gaa gat atc tgc aag ttt ttg aaa gca aaa act gga ttc cgt gtt cgc
I P Q L L E D I C K F F L L K A K T G G F R V R

cca gtc gcc gga tac tta tca gct cgt gat ttc ttg gca ggt ctt gca tat cgt gtc ttc
P V A G Y L S A R D F L L A G L A Y R R V F

ttc tgc act caa tac gtt cgc cat cat gcc gat cca ttt tac act cca gaa cca gac acc
F C T Q Y Y R R H H A D P F Y T P E P D T

gtt cac gag ctc atg ggt cac atg gct cta ttc gct gat cca gat ttt gct cag ttt tct
V H E L M G H A L F A D P F A Q F S

caa gag att gga tta gct tct ctt gga gca tca gag gaa gat ttg aag aag ctt gca aca
Q E I G L A S S L G A S E E D L K K L A T

ctc tac ttc ttt tcc att gaa ttt ggt ctc tcg tct gat gac gct gcc gat tct cca gta
L Y F F S I E F G L S S D D A A S P V

aaa gaa aat gga tca aat cat gaa aga ttt aaa gta tac gga gca ggt ctt ctg agc agt
K E N G S N H E R F F K V Y G A G L L S S

gct ggc gag ttg caa cat gcc gtt gag ggt agt gca acc att att cgt ttt gat ccg gat
A G E L Q H A V E G S A T I I R F D P D

cgt gtt gtt gag caa gaa tgt ctc att act act ttc cag tca gcg tat ttc tat act aga
R V V E Q E C L I T T F Q S A Y F Y T R

aat ttt gaa gag gcc cag cag aaa ctc aga atg ttc acc aac aac atg aaa cgt ccc ttc
N F E A Q Q K L R M F T N N M K R P F

att gtt cgt tac aac cca tac aca gaa agc gtc gaa gtt ctc aac aac tcc cgt tcc att
I V R Y N P Y T E S V L N N S R S I

atg ttg gca gtg aac tct ctc cgc tca gac atc aac ctc gct gga gct ctc cac tac
M L A V N S L R S D I N L L A G A L H Y

atc ctg tag
I L *

Fig. 42-2

attaccaccaagtttgaggtagcattgctctcttcaatcat

atg gat tgg ttg ttt cag atg gca tcc gca atg aag ttt caa tac tac tgg aag aaa gct
M D S L F Q M A S A M K F Q Y Y S K K A

gct gga aag aca atg tct aat agt gtc aaa aac tgg att ccg tgt tgg ccc agt cgc cgg
A G K T M S N S V K N W I P S R R

ata ctt atc agc tgc tga ttt ctt ggc agg tct tgc ata tgg tgt ctt ctt ctg cac tca
I L I S S *

ata cgt tgc cca tca tgc cga tcc att tta cac tcc aga acc aga cac cgt tca cga gct
cat ggg tca cat ggc tct att cgc tga tcc aga ttt tgc tca gtt ttc tca aga gat tgg
att agc ttc tct tgg agc atc aga gga aga ttt gaa gaa gct tgc aac act cta ctt ctt
ttc cat tga att tgg tct ctc gtc tga tga cgc tgc cga ttc tcc agt aaa aga aaa tgg
atc aaa tca tga aag att taa agt ata cgg agc agg act tct gag cag tgc tgg cga gtt
gca aca tgc cgt tga ggg tag tgc aac cat tat tgc ttt tga tcc gga tgc tgt tgt tga
gca aga atg tct cat tac ttt cca gtc agc gta ttt cta tac tag aaa ttt tga aga
ggc cca gca gaa act cag aat gtt cac caa cat gaa acg tcc ctt cat tgt tgc tta
caa ccc ata cac aga aag cgt cga agt tct caa cca ctc cgg ttc cat tat gtt ggc agt
gaa ctc tct ccg ctc aga cat caa cct gct cgc cgg agc tct cca cta cat cct gta g

Fig. 43

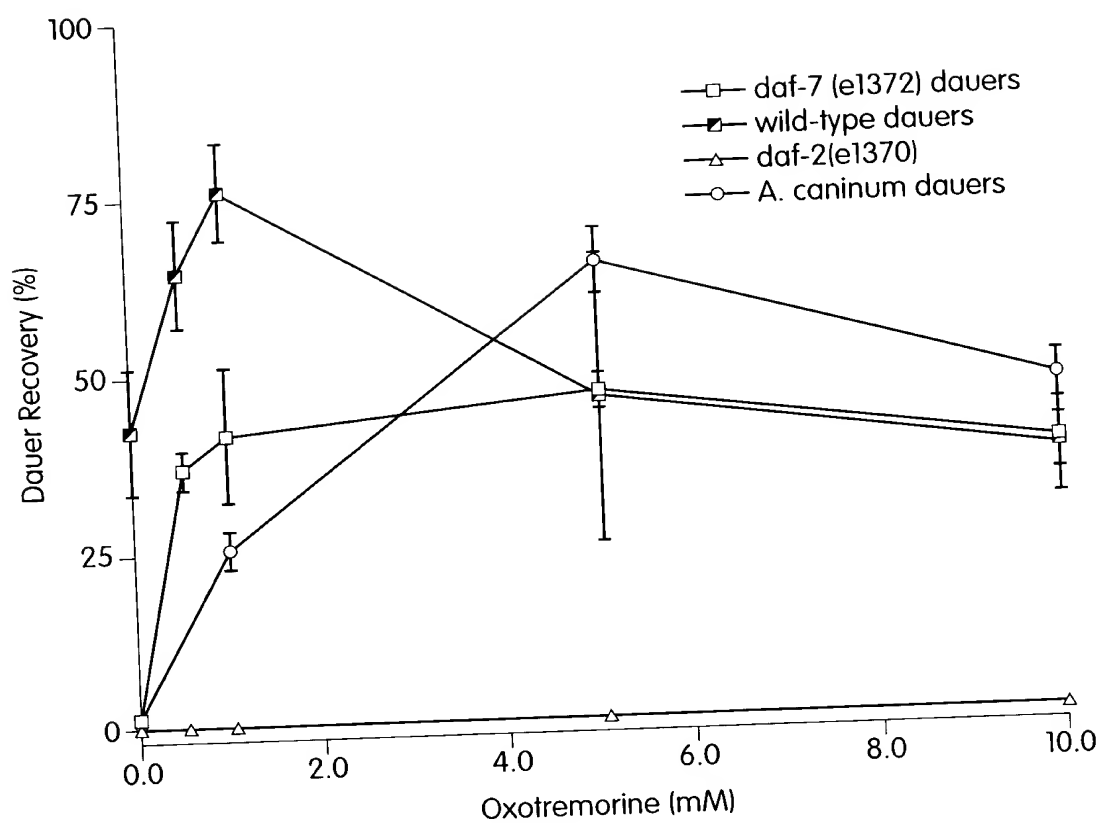


Fig. 44A

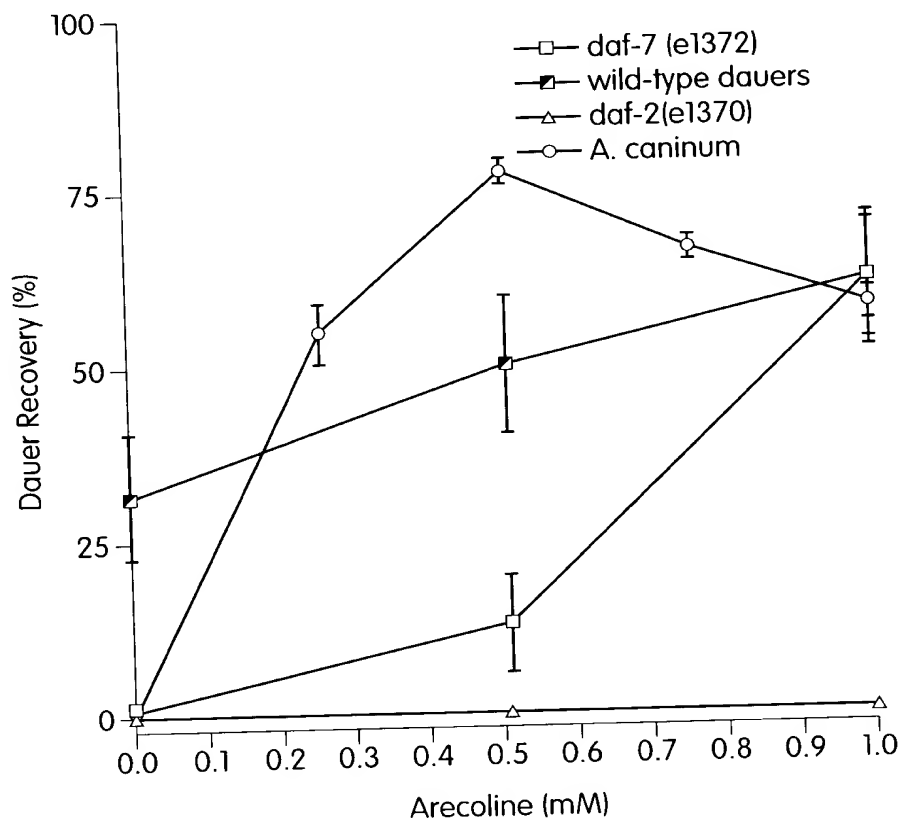


Fig. 44B

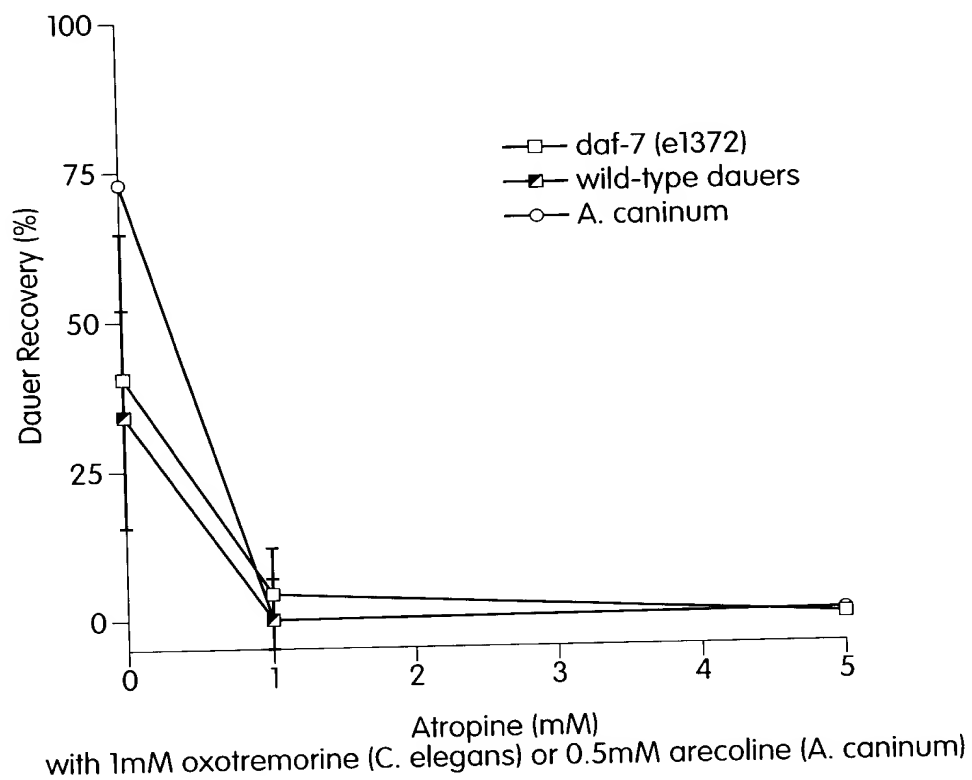


Fig. 44C



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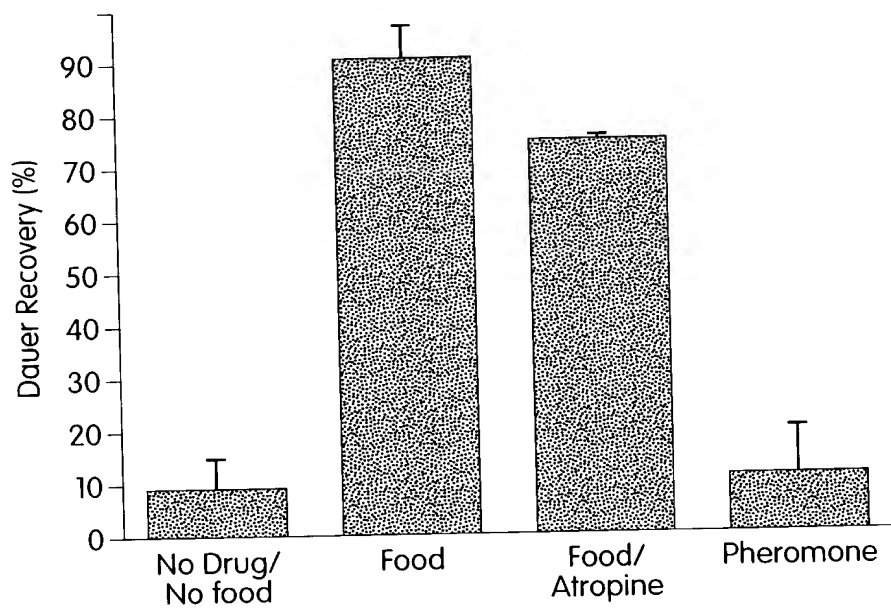


Fig. 45A



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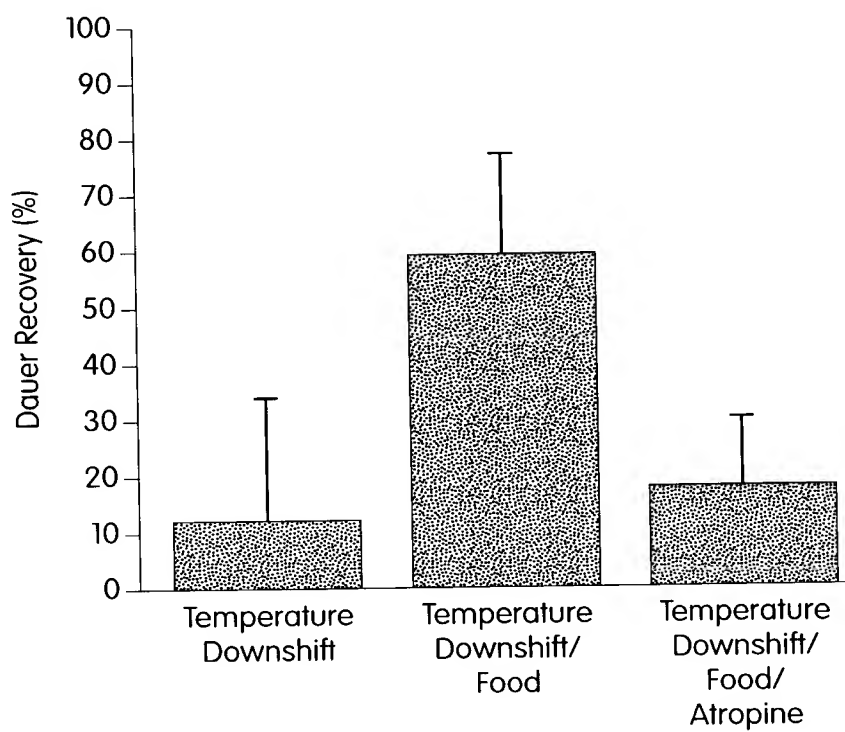


Fig. 45B

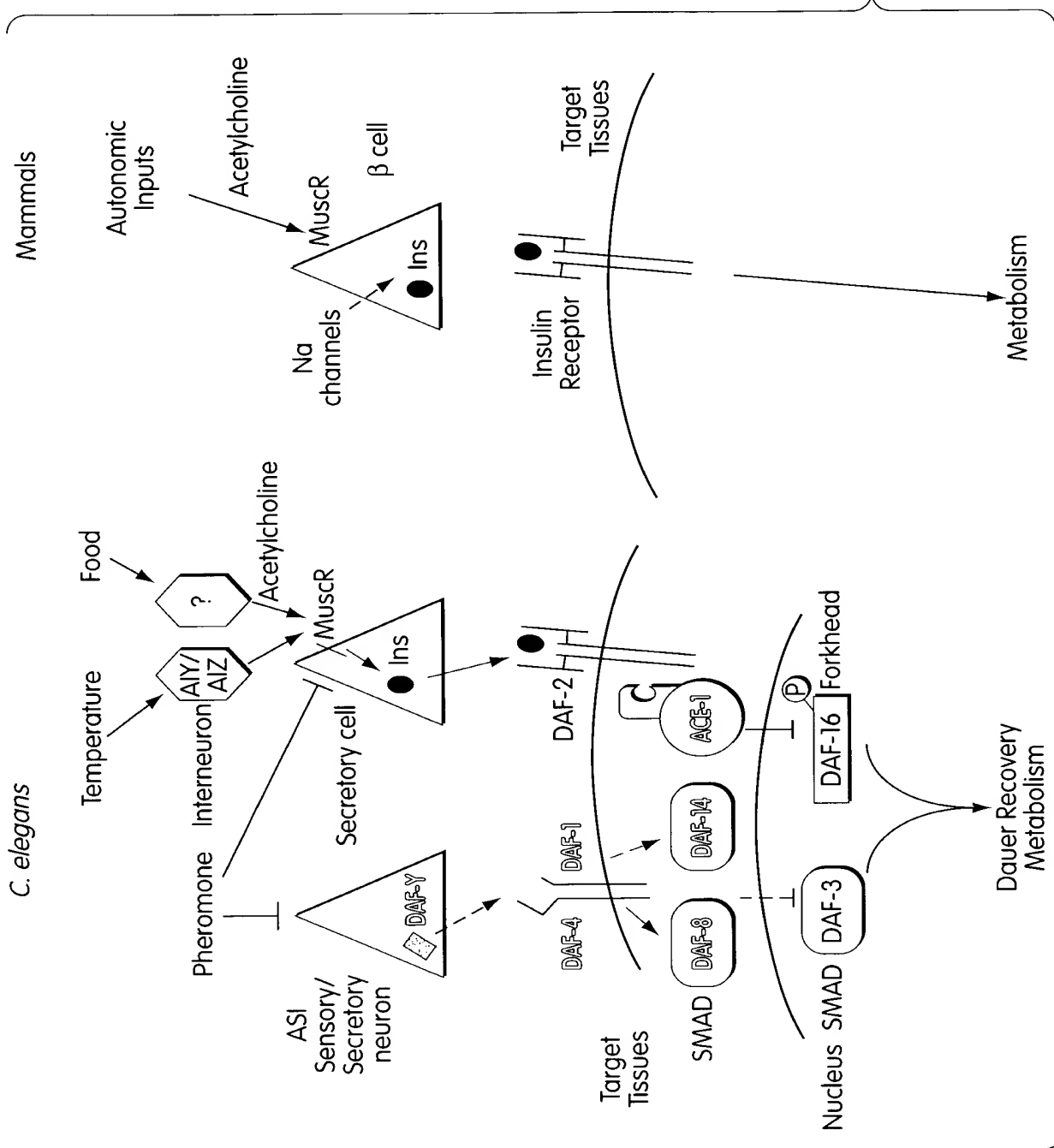


Fig. 46



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ATTCCGGCATGAGCATGGaGCTTCGAGTCCCTAGAGAACACAAACGTTCCCGCGGAACCTGGGtCTGGACTGGGAC
GAGACTCAAGCGAGTCCCGTCTGCTGCGGATATCCCTCACAGTGGACTTTGAGGCTTTGCGCTGGGACTGGATCAT
CGCACCTAAGCGCTACAAGGCCAACTACTGCTCCGGCCAGTGGAGTACATGTTTCATGCAAAAATATCCGCATACC
CATTTGGTGAGCAGGCCCAATCCAAAGAGTTATGCTGGGCCCTGTTGTACCCCCACCAAGATGTCCCAATCAACA
TgcTctACTTCAATGACAAAGCAGCAGATTTATctACGGCAAGATCCCTGGCATGTTGGTGGATCGCTGTGGctGCTC
TTAAGTGGGGATAGAGGATGCCTCCCCACAGACCGTACCCCCAAGACCCATAGCCctTGCCCAATCCACCGCCTG
ATCCAAACAT

Fig. 47A

IRHEHGASSPREHKTFPAEPGSGLRDRDSSSRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGQWEYMFQMYPHT
HLVQQANPRGYAGPCCTPTKMSPINMLYFNDKQIIYGKIPLAMVVDRCGCS

Fig. 47B